

SEQUENCE LISTING

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<110> Toray Industries Inc.

Tokyo Metropolitan Organization for Medical Research

Johannes Gutenberg-Universitaet Mainz

<120> A nucleic acid construct comprising genomic nucleic acid of genotype
2a Hepatitis C virus, and the nucleic acid construct-transferred cells

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<140> PCT/JP2003/015038

<141> 2003-11-25

<150> JP 2003-148242

<151> 2003-05-26

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<151> 2003-09-19

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<170> PatentIn Ver. 2.1

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<223> Inventor: Wakita, Takaji

Inventor: Kato, Takanobu

Inventor: Date, Tomoko

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Met Ser Thr Asn Pro

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5

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Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn Arg Arg Pro Glu Asp

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Pro	Arg	Arg	Gly	Pro	Arg	Leu	Gly	Val	Arg	Thr	Thr	Arg	Lys	Thr	Ser	
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Glu	Arg	Ser	Gln	Pro	Arg	Gly	Arg	Arg	Gln	Pro	Ile	Pro	Lys	Asp	Arg	
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Arg	Ser	Thr	Gly	Lys	Ala	Trp	Gly	Lys	Pro	Gly	Arg	Pro	Trp	Pro	Leu	
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Tyr	Gly	Asn	Glu	Gly	Leu	Gly	Trp	Ala	Gly	Trp	Leu	Leu	Ser	Pro	Arg	
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Gly	Ser	Arg	Pro	Ser	Trp	Gly	Pro	Thr	Asp	Pro	Arg	His	Arg	Ser	Arg	
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Asn	Val	Gly	Lys	Val	Ile	Asp	Thr	Leu	Thr	Cys	Gly	Phe	Ala	Asp	Leu	
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Leu Ser Cys Ile Thr Val Pro Val Ser Ala Ala Gln Val Lys Asn Thr			
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agt agc agc tac atg gtg acc aat gac tgc tcc aat gac agc atc act			979
Ser Ser Ser Tyr Met Val Thr Asn Asp Cys Ser Asn Asp Ser Ile Thr			
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Trp Gln Leu Glu Ala Ala Val Leu His Val Pro Gly Cys Val Pro Cys			
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280

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310

315

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325

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330

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360

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Asp Leu Leu Cys Pro Thr Asp Cys Phe Arg Lys His Pro Asp Ala Thr
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His Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe Thr			
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Ile Phe Lys Ile Arg Met Tyr Val Gly Gly Val Glu His Arg Leu Thr			
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gac agg agt cag ctg tct cct ctg ttg cac tct acc acg gaa tgg gcc			2371
Asp Arg Ser Gln Leu Ser Pro Leu Leu His Ser Thr Thr Glu Trp Ala			
665	670	675	
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Ile Leu Pro Cys Thr Tyr Ser Asp Leu Pro Ala Leu Ser Thr Gly Leu			
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Leu His Leu His Gln Asn Ile Val Asp Val Gln Tyr Met Tyr Gly Leu			
695	700	705	
tca cct gct atc aca aaa tac gtc gtt cga tgg gag tgg gtg gta ctc			2515
Ser Pro Ala Ile Thr Lys Tyr Val Val Arg Trp Glu Trp Val Val Leu			
710	715	720	725

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730

735

740

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745

750

755

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760

765

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775

780

785

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790

795

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Leu Ala Gly Leu Arg Gly Pro Val Thr Gln Met Tyr Ser Ser Ala Glu			
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Arg Thr Gly Arg Gly Arg Gln Gly Thr Tyr Arg Tyr Val Ser Thr Gly
1495 1500 1505

gaa cga gcc tca gga atg ttt gac agt gta gtg ctt tgt gag tgc tac 4915
Glu Arg Ala Ser Gly Met Phe Asp Ser Val Val Leu Cys Glu Cys Tyr
1510 1515 1520 1525

gac gca ggg gct gcg tgg tac gat ctc aca cca gcg gag acc acc gtc 4963
Asp Ala Gly Ala Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr Val

1530	1535	1540	
agg ctt aga gcg tat ttc aac acg ccc ggc cta ccc gtg tgt caa gac			5011
Arg Leu Arg Ala Tyr Phe Asn Thr Pro Gly Leu Pro Val Cys Gln Asp			
1545	1550	1555	
cat ctt gaa ttt tgg gag gca gtt ttc acc ggc ctc aca cac ata gac			5059
His Leu Glu Phe Trp Glu Ala Val Phe Thr Gly Leu Thr His Ile Asp			
1560	1565	1570	
gcc cac ttc ctc tcc caa aca aag caa gcg ggg gag aac ttc gcg tac			5107
Ala His Phe Leu Ser Gln Thr Lys Gln Ala Gly Glu Asn Phe Ala Tyr			
1575	1580	1585	
cta gta gcc tac caa gct acg gtg tgc gcc aga gcc aag gcc cct ccc			5155
Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Lys Ala Pro Pro			
1590	1595	1600	1605
ccg tcc tgg gac gcc atg tgg aag tgc ctg gcc cga ctc aag cct acg			5203
Pro Ser Trp Asp Ala Met Trp Lys Cys Leu Ala Arg Leu Lys Pro Thr			
1610	1615	1620	
ctt gcg ggc ccc aca cct ctc ctg tac cgt ttg ggc cct att acc aat			5251
Leu Ala Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Ile Thr Asn			
1625	1630	1635	
gag gtc acc ctc aca cac cct ggg acg aag tac atc gcc aca tgc atg			5299
Glu Val Thr Leu Thr His Pro Gly Thr Lys Tyr Ile Ala Thr Cys Met			
1640	1645	1650	

caa gct gac ctt gag gtc atg acc agc acg tgg gtc cta gct gga gga	5347
Gln Ala Asp Leu Glu Val Met Thr Ser Thr Trp Val Leu Ala Gly Gly	
1655 1660 1665	
gtc ctg gca gcc gtc gcc gca tat tgc ctg gcg act gga tgc gtt tcc	5395
Val Leu Ala Ala Val Ala Ala Tyr Cys Leu Ala Thr Gly Cys Val Ser	
1670 1675 1680 1685	
atc atc ggc cgc ttg cac gtc aac cag cga gtc gtc gtt gcg ccg gat	5443
Ile Ile Gly Arg Leu His Val Asn Gln Arg Val Val Val Ala Pro Asp	
1690 1695 1700	
aag gag gtc ctg tat gag gct ttt gat gag atg gag gaa tgc gcc tct	5491
Lys Glu Val Leu Tyr Glu Ala Phe Asp Glu Met Glu Glu Cys Ala Ser	
1705 1710 1715	
agg gcg gct ctc atc gaa gag ggg cag cgg ata gcc gag atg ttg aag	5539
Arg Ala Ala Leu Ile Glu Glu Gly Gln Arg Ile Ala Glu Met Leu Lys	
1720 1725 1730	
tcc aag atc caa ggc ttg ctg cag cag gcc tct aag cag gcc cag gac	5587
Ser Lys Ile Gln Gly Leu Leu Gln Gln Ala Ser Lys Gln Ala Gln Asp	
1735 1740 1745	
ata caa ccc gct atg cag gct tca tgg ccc aaa gtg gaa caa ttt tgg	5635
Ile Gln Pro Ala Met Gln Ala Ser Trp Pro Lys Val Glu Gln Phe Trp	
1750 1755 1760 1765	

gcc aga cac atg tgg aac ttc att agc ggc atc caa tac ctc gca gga 5683
Ala Arg His Met Trp Asn Phe Ile Ser Gly Ile Gln Tyr Leu Ala Gly
1770 1775 1780

ttg tca aca ctg cca ggg aac ccc gcg gtg gct tcc atg atg gca ttc 5731
Leu Ser Thr Leu Pro Gly Asn Pro Ala Val Ala Ser Met Met Ala Phe
1785 1790 1795

agt gcc gcc ctc acc agt ccg ttg tcg acc agt acc acc atc ctt ctc 5779
Ser Ala Ala Leu Thr Ser Pro Leu Ser Thr Ser Thr Thr Ile Leu Leu
1800 1805 1810

aac atc atg gga ggc tgg tta gcg tcc cag atc gca cca ccc gcg ggg 5827
Asn Ile Met Gly Gly Trp Leu Ala Ser Gln Ile Ala Pro Pro Ala Gly
1815 1820 1825

gcc acc ggc ttt gtc gtc agt ggc ctg gtg ggg gct gcc gtg ggc agc 5875
Ala Thr Gly Phe Val Val Ser Gly Leu Val Gly Ala Ala Val Gly Ser
1830 1835 1840 1845

ata ggc ctg ggt aag gtg ctg gtg gac atc ctg gca gga tat ggt gcg 5923
Ile Gly Leu Gly Lys Val Leu Val Asp Ile Leu Ala Gly Tyr Gly Ala
1850 1855 1860

ggc att tcg ggg gcc ctc gtc gca ttc aag atc atg tct ggc gag aag 5971
Gly Ile Ser Gly Ala Leu Val Ala Phe Lys Ile Met Ser Gly Glu Lys
1865 1870 1875

ccc tct atg gaa gat gtc atc aat cta ctg cct ggg atc ctg tct ccg 6019

Pro Ser Met Glu Asp Val Ile Asn Leu Leu Pro Gly Ile Leu Ser Pro
1880 1885 1890

gga gcc ctg gtg gtg ggg gtc atc tgc gcg gcc att ctg cgc cgc cac 6067
Gly Ala Leu Val Val Gly Val Ile Cys Ala Ala Ile Leu Arg Arg His
1895 1900 1905

gtg gga ccg ggg gag ggc gcg gtc caa tgg atg aac agg ctt att gcc 6115
Val Gly Pro Gly Glu Gly Ala Val Gln Trp Met Asn Arg Leu Ile Ala
1910 1915 1920 1925

ttt gct tcc aga gga aac cac gtc gcc cct act cac tac gtg acg gag 6163
Phe Ala Ser Arg Gly Asn His Val Ala Pro Thr His Tyr Val Thr Glu
1930 1935 1940

tcg gat gcg tcg cag cgt gtg acc caa cta ctt ggc tct ctt act ata 6211
Ser Asp Ala Ser Gln Arg Val Thr Gln Leu Leu Gly Ser Leu Thr Ile
1945 1950 1955

acc agc cta ctc aga aga ctc cac aat tgg ata act gag gac tgc ccc 6259
Thr Ser Leu Leu Arg Arg Leu His Asn Trp Ile Thr Glu Asp Cys Pro
1960 1965 1970

atc cca tgc tcc gga tcc tgg ctc cgc gac gtg tgg gac tgg gtt tgc 6307
Ile Pro Cys Ser Gly Ser Trp Leu Arg Asp Val Trp Asp Trp Val Cys
1975 1980 1985

acc atc ttg aca gac ttc aaa aat tgg ctg acc tct aaa ttg ttc ccc 6355
Thr Ile Leu Thr Asp Phe Lys Asn Trp Leu Thr Ser Lys Leu Phe Pro

1990	1995	2000	2005	
aag ctg ccc ggc ctc ccc ttc atc tct tgt caa aag ggg tac aag ggt				6403
Lys Leu Pro Gly Leu Pro Phe Ile Ser Cys Gln Lys Gly Tyr Lys Gly				
	2010	2015	2020	
gtg tgg gcc ggc act ggc atc atg acc acg cgc tgc cct tgc ggc gcc				6451
Val Trp Ala Gly Thr Gly Ile Met Thr Thr Arg Cys Pro Cys Gly Ala				
	2025	2030	2035	
aac atc tct ggc aat gtc cgc ctg ggc tct atg agg atc aca ggg cct				6499
Asn Ile Ser Gly Asn Val Arg Leu Gly Ser Met Arg Ile Thr Gly Pro				
	2040	2045	2050	
aaa acc tgc atg aac acc tgg cag ggg acc ttt cct atc aat tgc tac				6547
Lys Thr Cys Met Asn Thr Trp Gln Gly Thr Phe Pro Ile Asn Cys Tyr				
	2055	2060	2065	
acg gag ggc cag tgc gcg ccg aaa ccc ccc acg aac tac aag acc gcc				6595
Thr Glu Gly Gln Cys Ala Pro Lys Pro Pro Thr Asn Tyr Lys Thr Ala				
	2070	2075	2080	2085
atc tgg agg gtg gcg gcc tcg gag tac gcg gag gtg acg cag cat ggg				6643
Ile Trp Arg Val Ala Ala Ser Glu Tyr Ala Glu Val Thr Gln His Gly				
	2090	2095	2100	
tcg tac tcc tat gta aca gga ctg acc act gac aat ctg aaa att cct				6691
Ser Tyr Ser Tyr Val Thr Gly Leu Thr Thr Asp Asn Leu Lys Ile Pro				
	2105	2110	2115	

tgc caa cta cct tct cca gag ttt ttc tcc tgg gtg gac ggt gtg cag 6739
 Cys Gln Leu Pro Ser Pro Glu Phe Phe Ser Trp Val Asp Gly Val Gln

2120

2125

2130

atc cat agg ttt gca ccc aca cca aag ccg ttt ttc cgg gat gag gtc 6787
 Ile His Arg Phe Ala Pro Thr Pro Lys Pro Phe Phe Arg Asp Glu Val

2135

2140

2145

tcg ttc tgc gtt ggg ctt aat tcc tat gct gtc ggg tcc cag ctt ccc 6835
 Ser Phe Cys Val Gly Leu Asn Ser Tyr Ala Val Gly Ser Gln Leu Pro

2150

2155

2160

2165

tgt gaa cct gag ccc gac gca gac gta ttg agg tcc atg cta aca gat 6883
 Cys Glu Pro Glu Pro Asp Ala Asp Val Leu Arg Ser Met Leu Thr Asp

2170

2175

2180

ccg ccc cac atc acg gcg gag act gcg gcg cgg cgc ttg gca cgg gga 6931
 Pro Pro His Ile Thr Ala Glu Thr Ala Ala Arg Arg Leu Ala Arg Gly

2185

2190

2195

tca cct cca tct gag gcg agc tcc tca gtg agc cag cta tca gca ccg 6979
 Ser Pro Pro Ser Glu Ala Ser Ser Ser Val Ser Gln Leu Ser Ala Pro

2200

2205

2210

tcg ctg cgg gcc acc tgc acc acc cac agc aac acc tat gac gtg gac 7027
 Ser Leu Arg Ala Thr Cys Thr Thr His Ser Asn Thr Tyr Asp Val Asp

2215

2220

2225

atg gtc gat gcc aac ctg ctc atg gag ggc ggt gtg gct cag aca gag 7075
Met Val Asp Ala Asn Leu Leu Met Glu Gly Gly Val Ala Gln Thr Glu
2230 2235 2240 2245

cct gag tcc agg gtg ccc gtt ctg gac ttt ctc gag cca atg gcc gag 7123
Pro Glu Ser Arg Val Pro Val Leu Asp Phe Leu Glu Pro Met Ala Glu
2250 2255 2260

gaa gag agc gac ctt gag ccc tca ata cca tcg gag tgc atg ctc ccc 7171
Glu Glu Ser Asp Leu Glu Pro Ser Ile Pro Ser Glu Cys Met Leu Pro
2265 2270 2275

agg agc ggg ttt cca cgg gcc tta ccg gct tgg gca cgg cct gac tac 7219
Arg Ser Gly Phe Pro Arg Ala Leu Pro Ala Trp Ala Arg Pro Asp Tyr
2280 2285 2290

aac ccg ccg ctc gtg gaa tcg tgg agg agg cca gat tac caa ccg ccc 7267
Asn Pro Pro Leu Val Glu Ser Trp Arg Arg Pro Asp Tyr Gln Pro Pro
2295 2300 2305

acc gtt gct ggt tgt gct ctc ccc ccc ccc aag aag gcc ccg acg cct 7315
Thr Val Ala Gly Cys Ala Leu Pro Pro Pro Lys Lys Ala Pro Thr Pro
2310 2315 2320 2325

ccc cca agg aga cgc cgg aca gtg ggt ctg agc gag agc acc ata tca 7363
Pro Pro Arg Arg Arg Arg Thr Val Gly Leu Ser Glu Ser Thr Ile Ser
2330 2335 2340

gaa gcc ctc cag caa ctg gcc atc aag acc ttt ggc cag ccc ccc tcg 7411

Glu Ala Leu Gln Gln Leu Ala Ile Lys Thr Phe Gly Gln Pro Pro Ser	
2345	2350 2355
agc ggt gat gca ggc tcg tcc acg ggg gcg ggc gcc gcc gaa tcc ggc	7459
Ser Gly Asp Ala Gly Ser Ser Thr Gly Ala Gly Ala Ala Glu Ser Gly	
2360	2365 2370
ggt ccg acg tcc cct ggt gag ccg gcc ccc tca gag aca ggt tcc gcc	7507
Gly Pro Thr Ser Pro Gly Glu Pro Ala Pro Ser Glu Thr Gly Ser Ala	
2375	2380 2385
tcc tct atg ccc ccc ctc gag ggg gag cct gga gat ccg gac ctg gag	7555
Ser Ser Met Pro Pro Leu Glu Gly Glu Pro Gly Asp Pro Asp Leu Glu	
2390	2395 2400 2405
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Ser Asp Gln Val Glu Leu Gln Pro Pro Pro Gln Gly Gly Gly Val Ala	
2410	2415 2420
ccc ggt tcg ggc tcg ggg tct tgg tct act tgc tcc gag gag gac gat	7651
Pro Gly Ser Gly Ser Gly Ser Trp Ser Thr Cys Ser Glu Glu Asp Asp	
2425	2430 2435
acc acc gtg tgc tgc tcc atg tca tac tcc tgg acc ggg gct cta ata	7699
Thr Thr Val Cys Cys Ser Met Ser Tyr Ser Trp Thr Gly Ala Leu Ile	
2440	2445 2450
act ccc tgt agc ccc gaa gag gaa aag ttg cca atc aac cct ttg agt	7747
Thr Pro Cys Ser Pro Glu Glu Glu Lys Leu Pro Ile Asn Pro Leu Ser	

2455	2460	2465	
aac tcg ctg ttg cga tac cat aac aag gtg tac tgt aca aca tca aag			7795
Asn Ser Leu Leu Arg Tyr His Asn Lys Val Tyr Cys Thr Thr Ser Lys			
2470	2475	2480	2485
agc gcc tca cag agg gct aaa aag gta act ttt gac agg acg caa gtg			7843
Ser Ala Ser Gln Arg Ala Lys Lys Val Thr Phe Asp Arg Thr Gln Val			
	2490	2495	2500
ctc gac gcc cat tat gac tca gtc tta aag gac atc aag cta gcg gct			7891
Leu Asp Ala His Tyr Asp Ser Val Leu Lys Asp Ile Lys Leu Ala Ala			
	2505	2510	2515
tcc aag gtc agc gca agg ctc ctc acc ttg gag gag gcg tgc cag ttg			7939
Ser Lys Val Ser Ala Arg Leu Leu Thr Leu Glu Glu Ala Cys Gln Leu			
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act cca ccc cat tct gca aga tcc aag tat gga ttc ggg gcc aag gag			7987
Thr Pro Pro His Ser Ala Arg Ser Lys Tyr Gly Phe Gly Ala Lys Glu			
	2535	2540	2545
gtc cgc agc ttg tcc ggg agg gcc gtt aac cac atc aag tcc gtg tgg			8035
Val Arg Ser Leu Ser Gly Arg Ala Val Asn His Ile Lys Ser Val Trp			
2550	2555	2560	2565
aag gac ctc ctg gaa gac cca caa aca cca att ccc aca acc atc atg			8083
Lys Asp Leu Leu Glu Asp Pro Gln Thr Pro Ile Pro Thr Thr Ile Met			
	2570	2575	2580

gcc aaa aat gag gtg ttc tgc gtg gac ccc gcc aag ggg ggt aag aaa 8131
Ala Lys Asn Glu Val Phe Cys Val Asp Pro Ala Lys Gly Gly Lys Lys

2585

2590

2595

cca gct cgc ctc atc gtt tac cct gac ctc ggc gtc cgg gtc tgc gag 8179
Pro Ala Arg Leu Ile Val Tyr Pro Asp Leu Gly Val Arg Val Cys Glu

2600

2605

2610

aaa atg gcc ctc tat gac att aca caa aag ctt cct cag gcg gta atg 8227
Lys Met Ala Leu Tyr Asp Ile Thr Gln Lys Leu Pro Gln Ala Val Met

2615

2620

2625

gga gct tcc tat ggc ttc cag tac tcc cct gcc caa cgg gtg gag tat 8275
Gly Ala Ser Tyr Gly Phe Gln Tyr Ser Pro Ala Gln Arg Val Glu Tyr

2630

2635

2640

2645

ctc ttg aaa gca tgg gcg gaa aag aag gac ccc atg ggt ttt tcg tat 8323
Leu Leu Lys Ala Trp Ala Glu Lys Lys Asp Pro Met Gly Phe Ser Tyr

2650

2655

2660

gat acc cga tgc ttc gac tca acc gtc act gag aga gac atc agg acc 8371
Asp Thr Arg Cys Phe Asp Ser Thr Val Thr Glu Arg Asp Ile Arg Thr

2665

2670

2675

gag gag tcc ata tac cag gcc tgc tcc ctg ccc gag gag gcc cgc act 8419
Glu Glu Ser Ile Tyr Gln Ala Cys Ser Leu Pro Glu Glu Ala Arg Thr

2680

2685

2690

gcc ata cac tcg ctg act gag aga ctt tac gta gga ggg ccc atg ttc 8467
 Ala Ile His Ser Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Met Phe
 2695 2700 2705

aac agc aag ggt caa acc tgc ggt tac aga cgt tgc cgc gcc agc ggg 8515
 Asn Ser Lys Gly Gln Thr Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly
 2710 2715 2720 2725

gtg cta acc act agc atg ggt aac acc atc aca tgc tat gtg aaa gcc 8563
 Val Leu Thr Thr Ser Met Gly Asn Thr Ile Thr Cys Tyr Val Lys Ala
 2730 2735 2740

cta gcg gcc tgc aag gct gcg ggg ata gtt gcg ccc aca atg ctg gta 8611
 Leu Ala Ala Cys Lys Ala Ala Gly Ile Val Ala Pro Thr Met Leu Val
 2745 2750 2755

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 Cys Gly Asp Asp Leu Val Val Ile Ser Glu Ser Gln Gly Thr Glu Glu
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 Asp Glu Arg Asn Leu Arg Ala Phe Thr Glu Ala Met Thr Arg Tyr Ser
 2775 2780 2785

gcc cct cct ggt gat ccc ccc aga ccg gaa tat gac ctg gag cta ata 8755
 Ala Pro Pro Gly Asp Pro Pro Arg Pro Glu Tyr Asp Leu Glu Leu Ile
 2790 2795 2800 2805

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Thr Ser Cys Ser Ser Asn Val Ser Val Ala Leu Gly Pro Arg Gly Arg	
2810	2815 2820
cgc aga tac tac ctg acc aga gac cca acc act cca ctc gcc cgg gct	8851
Arg Arg Tyr Tyr Leu Thr Arg Asp Pro Thr Thr Pro Leu Ala Arg Ala	
2825	2830 2835
gcc tgg gaa aca gtt aga cac tcc cct atc aat tca tgg ctg gga aac	8899
Ala Trp Glu Thr Val Arg His Ser Pro Ile Asn Ser Trp Leu Gly Asn	
2840	2845 2850
atc atc cag tat gct cca acc ata tgg gtt cgc atg gtc cta atg aca	8947
Ile Ile Gln Tyr Ala Pro Thr Ile Trp Val Arg Met Val Leu Met Thr	
2855	2860 2865
cac ttc ttc tcc att ctc atg gtc caa gac acc ctg gac cag aac ctc	8995
His Phe Phe Ser Ile Leu Met Val Gln Asp Thr Leu Asp Gln Asn Leu	
2870	2875 2880 2885
aac ttt gag atg tat gga tca gta tac tcc gtg aat cct ttg gac ctt	9043
Asn Phe Glu Met Tyr Gly Ser Val Tyr Ser Val Asn Pro Leu Asp Leu	
2890	2895 2900
cca gcc ata att gag agg tta cac ggg ctt gac gcc ttt tct atg cac	9091
Pro Ala Ile Ile Glu Arg Leu His Gly Leu Asp Ala Phe Ser Met His	
2905	2910 2915
aca tac tct cac cac gaa ctg acg cgg gtg gct tca gcc ctc aga aaa	9139
Thr Tyr Ser His His Glu Leu Thr Arg Val Ala Ser Ala Leu Arg Lys	

2920	2925	2930	
ctt ggg gcg cca ccc ctc agg gtg tgg aag agt cgg gct cgc gca gtc			9187
Leu Gly Ala Pro Pro Leu Arg Val Trp Lys Ser Arg Ala Arg Ala Val			
2935	2940	2945	
agg gcg tcc ctc atc tcc cgt gga ggg aaa gcg gcc gtt tgc ggc cga			9235
Arg Ala Ser Leu Ile Ser Arg Gly Gly Lys Ala Ala Val Cys Gly Arg			
2950	2955	2960	2965
tat ctc ttc aat tgg gcg gtg aag acc aag ctc aaa ctc act cca ttg			9283
Tyr Leu Phe Asn Trp Ala Val Lys Thr Lys Leu Lys Leu Thr Pro Leu			
2970	2975	2980	
ccg gag gcg cgc cta ctg gac tta tcc agt tgg ttc acc gtc ggc gcc			9331
Pro Glu Ala Arg Leu Leu Asp Leu Ser Ser Trp Phe Thr Val Gly Ala			
2985	2990	2995	
ggc ggg ggc gac att ttt cac agc gtg tcg cgc gcc cga ccc cgc tca			9379
Gly Gly Gly Asp Ile Phe His Ser Val Ser Arg Ala Arg Pro Arg Ser			
3000	3005	3010	
tta ctc ttc ggc cta ctc cta ctt ttc gta ggg gta ggc ctc ttc cta			9427
Leu Leu Phe Gly Leu Leu Leu Leu Phe Val Gly Val Gly Leu Phe Leu			
3015	3020	3025	
ctc ccc gct cgg tag agcggcacac actaggtaca ctccatagct aactgttcct			9482
Leu Pro Ala Arg			
3030			

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tctctgcaga tcatgt 9678

<210> 4

<211> 3033

<212> PRT

<213> Hepatitis C virus

<400> 4

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn

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Arg Arg Pro Glu Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly

20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Thr

35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro

50 55 60

Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ala Trp Gly Lys Pro Gly

65 70 75 80

Arg Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp

85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro

100	105	110
Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys		
115	120	125
Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Leu		
130	135	140
Ser Gly Ala Ala Arg Ala Val Ala His Gly Val Arg Val Leu Glu Asp		
145	150	155
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Phe Pro Phe Ser Ile		
165	170	175
Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Val Pro Val Ser Ala Ala		
180	185	190
Gln Val Lys Asn Thr Ser Ser Ser Tyr Met Val Thr Asn Asp Cys Ser		
195	200	205
Asn Asp Ser Ile Thr Trp Gln Leu Glu Ala Ala Val Leu His Val Pro		
210	215	220
Gly Cys Val Pro Cys Glu Arg Val Gly Asn Thr Ser Arg Cys Trp Val		
225	230	235
Pro Val Ser Pro Asn Met Ala Val Arg Gln Pro Gly Ala Leu Thr Gln		
245	250	255
Gly Leu Arg Thr His Ile Asp Met Val Val Met Ser Ala Thr Phe Cys		
260	265	270
Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Val Met Leu Ala Ala		
275	280	285
Gln Val Phe Ile Val Ser Pro Gln Tyr His Trp Phe Val Gln Glu Cys		
290	295	300
Asn Cys Ser Ile Tyr Pro Gly Thr Ile Thr Gly His Arg Met Ala Trp		
305	310	315
Asp Met Met Met Asn Trp Ser Pro Thr Ala Thr Met Ile Leu Ala Tyr		
325	330	335

Val Met Arg Val Pro Glu Val Ile Ile Asp Ile Val Ser Gly Ala His
340 345 350
Trp Gly Val Met Phe Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala Trp
355 360 365
Ala Lys Val Ile Val Ile Leu Leu Leu Ala Ala Gly Val Asp Ala Gly
370 375 380
Thr Thr Thr Val Gly Gly Ala Val Ala Arg Ser Thr Asn Val Ile Ala
385 390 395 400
Gly Val Phe Ser His Gly Pro Gln Gln Asn Ile Gln Leu Ile Asn Thr
405 410 415
Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser
420 425 430
Leu Asn Thr Gly Phe Leu Ala Ala Leu Phe Tyr Thr Asn Arg Phe Asn
435 440 445
Ser Ser Gly Cys Pro Gly Arg Leu Ser Ala Cys Arg Asn Ile Glu Ala
450 455 460
Phe Arg Ile Gly Trp Gly Thr Leu Gln Tyr Glu Asp Asn Val Thr Asn
465 470 475 480
Pro Glu Asp Met Arg Pro Tyr Cys Trp His Tyr Pro Pro Lys Pro Cys
485 490 495
Gly Val Val Pro Ala Arg Ser Val Cys Gly Pro Val Tyr Cys Phe Thr
500 505 510
Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Arg Gly Val Pro Thr
515 520 525
Tyr Thr Trp Gly Glu Asn Glu Thr Asp Val Phe Leu Leu Asn Ser Thr
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Arg Pro Pro Gln Gly Ser Trp Phe Gly Cys Thr Trp Met Asn Ser Thr
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Gly Phe Thr Lys Thr Cys Gly Ala Pro Pro Cys Arg Thr Arg Ala Asp

	565		570		575										
Phe	Asn	Ala	Ser	Thr	Asp	Leu	Leu	Cys	Pro	Thr	Asp	Cys	Phe	Arg	Lys
	580						585						590		
His	Pro	Asp	Ala	Thr	Tyr	Ile	Lys	Cys	Gly	Ser	Gly	Pro	Trp	Leu	Thr
	595						600						605		
Pro	Lys	Cys	Leu	Val	His	Tyr	Pro	Tyr	Arg	Leu	Trp	His	Tyr	Pro	Cys
	610						615						620		
Thr	Val	Asn	Phe	Thr	Ile	Phe	Lys	Ile	Arg	Met	Tyr	Val	Gly	Gly	Val
625							630				635				640
Glu	His	Arg	Leu	Thr	Ala	Ala	Cys	Asn	Phe	Thr	Arg	Gly	Asp	Arg	Cys
							645				650				655
Asp	Leu	Glu	Asp	Arg	Asp	Arg	Ser	Gln	Leu	Ser	Pro	Leu	Leu	His	Ser
	660							665						670	
Thr	Thr	Glu	Trp	Ala	Ile	Leu	Pro	Cys	Thr	Tyr	Ser	Asp	Leu	Pro	Ala
	675							680						685	
Leu	Ser	Thr	Gly	Leu	Leu	His	Leu	His	Gln	Asn	Ile	Val	Asp	Val	Gln
	690							695						700	
Tyr	Met	Tyr	Gly	Leu	Ser	Pro	Ala	Ile	Thr	Lys	Tyr	Val	Val	Arg	Trp
705							710				715				720
Glu	Trp	Val	Val	Leu	Leu	Phe	Leu	Leu	Leu	Ala	Asp	Ala	Arg	Val	Cys
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Ala	Cys	Leu	Trp	Met	Leu	Ile	Leu	Leu	Gly	Gln	Ala	Glu	Ala	Ala	Leu
	740							745						750	
Glu	Lys	Leu	Val	Val	Leu	His	Ala	Ala	Ser	Ala	Ala	Asn	Cys	His	Gly
	755							760						765	
Leu	Leu	Tyr	Phe	Ala	Ile	Phe	Phe	Val	Ala	Ala	Trp	His	Ile	Arg	Gly
	770							775						780	
Arg	Val	Val	Pro	Leu	Thr	Thr	Tyr	Cys	Leu	Thr	Gly	Leu	Trp	Pro	Phe
785							790				795				800

Cys	Leu	Leu	Leu	Met	Ala	Leu	Pro	Arg	Gln	Ala	Tyr	Ala	Tyr	Asp	Ala
				805					810					815	
Pro	Val	His	Gly	Gln	Ile	Gly	Val	Gly	Leu	Leu	Ile	Leu	Ile	Thr	Leu
			820					825					830		
Phe	Thr	Leu	Thr	Pro	Gly	Tyr	Lys	Thr	Leu	Leu	Gly	Gln	Cys	Leu	Trp
		835					840					845			
Trp	Leu	Cys	Tyr	Leu	Leu	Thr	Leu	Gly	Glu	Ala	Met	Ile	Gln	Glu	Trp
	850					855					860				
Val	Pro	Pro	Met	Gln	Val	Arg	Gly	Gly	Arg	Asp	Gly	Ile	Ala	Trp	Ala
865					870					875				880	
Val	Thr	Ile	Phe	Cys	Pro	Gly	Val	Val	Phe	Asp	Ile	Thr	Lys	Trp	Leu
			885					890					895		
Leu	Ala	Leu	Leu	Gly	Pro	Ala	Tyr	Leu	Leu	Arg	Ala	Ala	Leu	Thr	His
		900					905					910			
Val	Pro	Tyr	Phe	Val	Arg	Ala	His	Ala	Leu	Ile	Arg	Val	Cys	Ala	Leu
	915						920				925				
Val	Lys	Gln	Leu	Ala	Gly	Gly	Arg	Tyr	Val	Gln	Val	Ala	Leu	Leu	Ala
	930				935					940					
Leu	Gly	Arg	Trp	Thr	Gly	Thr	Tyr	Ile	Tyr	Asp	His	Leu	Thr	Pro	Met
945					950					955				960	
Ser	Asp	Trp	Ala	Ala	Ser	Gly	Leu	Arg	Asp	Leu	Ala	Val	Ala	Val	Glu
			965					970				975			
Pro	Ile	Ile	Phe	Ser	Pro	Met	Glu	Lys	Lys	Val	Ile	Val	Trp	Gly	Ala
		980						985				990			
Glu	Thr	Ala	Ala	Cys	Gly	Asp	Ile	Leu	His	Gly	Leu	Pro	Val	Ser	Ala
	995					1000				1005					
Arg	Leu	Gly	Gln	Glu	Ile	Leu	Leu	Gly	Pro	Ala	Asp	Gly	Tyr	Thr	Ser
1010					1015					1020					
Lys	Gly	Trp	Lys	Leu	Leu	Ala	Pro	Ile	Thr	Ala	Tyr	Ala	Gln	Gln	Thr

1025	1030	1035	1040
Arg Gly Leu Leu Gly Ala Ile Val Val Ser Met Thr Gly Arg Asp Arg			
	1045	1050	1055
Thr Glu Gln Ala Gly Glu Val Gln Ile Leu Ser Thr Val Ser Gln Ser			
	1060	1065	1070
Phe Leu Gly Thr Thr Ile Ser Gly Val Leu Trp Thr Val Tyr His Gly			
	1075	1080	1085
Ala Gly Asn Lys Thr Leu Ala Gly Leu Arg Gly Pro Val Thr Gln Met			
	1090	1095	1100
Tyr Ser Ser Ala Glu Gly Asp Leu Val Gly Trp Pro Ser Pro Pro Gly			
1105	1110	1115	1120
Thr Lys Ser Leu Glu Pro Cys Lys Cys Gly Ala Val Asp Leu Tyr Leu			
	1125	1130	1135
Val Thr Arg Asn Ala Asp Val Ile Pro Ala Arg Arg Arg Gly Asp Lys			
	1140	1145	1150
Arg Gly Ala Leu Leu Ser Pro Arg Pro Ile Ser Thr Leu Lys Gly Ser			
	1155	1160	1165
Ser Gly Gly Pro Val Leu Cys Pro Arg Gly His Val Val Gly Leu Phe			
	1170	1175	1180
Arg Ala Ala Val Cys Ser Arg Gly Val Ala Lys Ser Ile Asp Phe Ile			
1185	1190	1195	1200
Pro Val Glu Thr Leu Asp Val Val Thr Arg Ser Pro Thr Phe Ser Asp			
	1205	1210	1215
Asn Ser Thr Pro Pro Ala Val Pro Gln Thr Tyr Gln Val Gly Tyr Leu			
	1220	1225	1230
His Ala Pro Thr Gly Ser Gly Lys Ser Thr Lys Val Pro Val Ala Tyr			
	1235	1240	1245
Ala Ala Gln Gly Tyr Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala			
	1250	1255	1260

Thr Leu Gly Phe Gly Ala Tyr Leu Ser Lys Ala His Gly Ile Asn Pro			
1265	1270	1275	1280
Asn Ile Arg Thr Gly Val Arg Thr Val Met Thr Gly Glu Ala Ile Thr			
	1285	1290	1295
Tyr Ser Thr Tyr Gly Lys Phe Leu Ala Asp Gly Gly Cys Ala Ser Gly			
1300	1305	1310	
Ala Tyr Asp Ile Ile Ile Cys Asp Glu Cys His Ala Val Asp Ala Thr			
1315	1320	1325	
Ser Ile Leu Gly Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly			
1330	1335	1340	
Val Arg Leu Thr Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr			
1345	1350	1355	1360
Thr Pro His Pro Asp Ile Glu Glu Val Gly Leu Gly Arg Glu Gly Glu			
	1365	1370	1375
Ile Pro Phe Tyr Gly Arg Ala Ile Pro Leu Ser Cys Ile Lys Gly Gly			
1380	1385	1390	
Arg His Leu Ile Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala			
1395	1400	1405	
Ala Ala Leu Arg Gly Met Gly Leu Asn Ala Val Ala Tyr Tyr Arg Gly			
1410	1415	1420	
Leu Asp Val Ser Ile Ile Pro Ala Gln Gly Asp Val Val Val Val Ala			
1425	1430	1435	1440
Thr Asp Ala Leu Met Thr Gly Tyr Thr Gly Asp Phe Asp Ser Val Ile			
	1445	1450	1455
Asp Cys Asn Val Ala Val Thr Gln Ala Val Asp Phe Ser Leu Asp Pro			
1460	1465	1470	
Thr Phe Thr Ile Thr Thr Gln Thr Val Pro Gln Asp Ala Val Ser Arg			
1475	1480	1485	
Ser Gln Arg Arg Gly Arg Thr Gly Arg Gly Arg Gln Gly Thr Tyr Arg			

1490	1495	1500
Tyr Val Ser Thr Gly Glu Arg Ala Ser Gly Met Phe Asp Ser Val Val		
1505	1510	1515
Leu Cys Glu Cys Tyr Asp Ala Gly Ala Ala Trp Tyr Asp Leu Thr Pro		1520
	1525	1530
Ala Glu Thr Thr Val Arg Leu Arg Ala Tyr Phe Asn Thr Pro Gly Leu		1535
	1540	1545
Pro Val Cys Gln Asp His Leu Glu Phe Trp Glu Ala Val Phe Thr Gly		1550
	1555	1560
Leu Thr His Ile Asp Ala His Phe Leu Ser Gln Thr Lys Gln Ala Gly		1565
	1570	1575
Glu Asn Phe Ala Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg		1580
1585	1590	1595
Ala Lys Ala Pro Pro Pro Ser Trp Asp Ala Met Trp Lys Cys Leu Ala		1600
	1605	1610
Arg Leu Lys Pro Thr Leu Ala Gly Pro Thr Pro Leu Leu Tyr Arg Leu		1615
	1620	1625
Gly Pro Ile Thr Asn Glu Val Thr Leu Thr His Pro Gly Thr Lys Tyr		1630
	1635	1640
Ile Ala Thr Cys Met Gln Ala Asp Leu Glu Val Met Thr Ser Thr Trp		1645
	1650	1655
Val Leu Ala Gly Gly Val Leu Ala Ala Val Ala Ala Tyr Cys Leu Ala		1660
1665	1670	1675
Thr Gly Cys Val Ser Ile Ile Gly Arg Leu His Val Asn Gln Arg Val		1680
	1685	1690
Val Val Ala Pro Asp Lys Glu Val Leu Tyr Glu Ala Phe Asp Glu Met		1695
	1700	1705
Glu Glu Cys Ala Ser Arg Ala Ala Leu Ile Glu Glu Gly Gln Arg Ile		1710
	1715	1720
		1725

Ala Glu Met Leu Lys Ser Lys Ile Gln Gly Leu Leu Gln Gln Ala Ser
 1730 1735 1740
 Lys Gln Ala Gln Asp Ile Gln Pro Ala Met Gln Ala Ser Trp Pro Lys
 1745 1750 1755 1760
 Val Glu Gln Phe Trp Ala Arg His Met Trp Asn Phe Ile Ser Gly Ile
 1765 1770 1775
 Gln Tyr Leu Ala Gly Leu Ser Thr Leu Pro Gly Asn Pro Ala Val Ala
 1780 1785 1790
 Ser Met Met Ala Phe Ser Ala Ala Leu Thr Ser Pro Leu Ser Thr Ser
 1795 1800 1805
 Thr Thr Ile Leu Leu Asn Ile Met Gly Gly Trp Leu Ala Ser Gln Ile
 1810 1815 1820
 Ala Pro Pro Ala Gly Ala Thr Gly Phe Val Val Ser Gly Leu Val Gly
 1825 1830 1835 1840
 Ala Ala Val Gly Ser Ile Gly Leu Gly Lys Val Leu Val Asp Ile Leu
 1845 1850 1855
 Ala Gly Tyr Gly Ala Gly Ile Ser Gly Ala Leu Val Ala Phe Lys Ile
 1860 1865 1870
 Met Ser Gly Glu Lys Pro Ser Met Glu Asp Val Ile Asn Leu Leu Pro
 1875 1880 1885
 Gly Ile Leu Ser Pro Gly Ala Leu Val Val Gly Val Ile Cys Ala Ala
 1890 1895 1900
 Ile Leu Arg Arg His Val Gly Pro Gly Glu Gly Ala Val Gln Trp Met
 1905 1910 1915 1920
 Asn Arg Leu Ile Ala Phe Ala Ser Arg Gly Asn His Val Ala Pro Thr
 1925 1930 1935
 His Tyr Val Thr Glu Ser Asp Ala Ser Gln Arg Val Thr Gln Leu Leu
 1940 1945 1950
 Gly Ser Leu Thr Ile Thr Ser Leu Leu Arg Arg Leu His Asn Trp Ile

1955	1960	1965
Thr Glu Asp Cys Pro Ile Pro Cys Ser Gly Ser Trp Leu Arg Asp Val		
1970	1975	1980
Trp Asp Trp Val Cys Thr Ile Leu Thr Asp Phe Lys Asn Trp Leu Thr		
1985	1990	1995
Ser Lys Leu Phe Pro Lys Leu Pro Gly Leu Pro Phe Ile Ser Cys Gln		2000
2005	2010	2015
Lys Gly Tyr Lys Gly Val Trp Ala Gly Thr Gly Ile Met Thr Thr Arg		
2020	2025	2030
Cys Pro Cys Gly Ala Asn Ile Ser Gly Asn Val Arg Leu Gly Ser Met		
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Arg Ile Thr Gly Pro Lys Thr Cys Met Asn Thr Trp Gln Gly Thr Phe		
2050	2055	2060
Pro Ile Asn Cys Tyr Thr Glu Gly Gln Cys Ala Pro Lys Pro Pro Thr		
2065	2070	2075
Asn Tyr Lys Thr Ala Ile Trp Arg Val Ala Ala Ser Glu Tyr Ala Glu		2080
2085	2090	2095
Val Thr Gln His Gly Ser Tyr Ser Tyr Val Thr Gly Leu Thr Thr Asp		
2100	2105	2110
Asn Leu Lys Ile Pro Cys Gln Leu Pro Ser Pro Glu Phe Phe Ser Trp		
2115	2120	2125
Val Asp Gly Val Gln Ile His Arg Phe Ala Pro Thr Pro Lys Pro Phe		
2130	2135	2140
Phe Arg Asp Glu Val Ser Phe Cys Val Gly Leu Asn Ser Tyr Ala Val		
2145	2150	2155
Gly Ser Gln Leu Pro Cys Glu Pro Glu Pro Asp Ala Asp Val Leu Arg		2160
2165	2170	2175
Ser Met Leu Thr Asp Pro Pro His Ile Thr Ala Glu Thr Ala Ala Arg		
2180	2185	2190

Arg Leu Ala Arg Gly Ser Pro Pro Ser Glu Ala Ser Ser Ser Val Ser
 2195 2200 2205
 Gln Leu Ser Ala Pro Ser Leu Arg Ala Thr Cys Thr Thr His Ser Asn
 2210 2215 2220
 Thr Tyr Asp Val Asp Met Val Asp Ala Asn Leu Leu Met Glu Gly Gly
 2225 2230 2235 2240
 Val Ala Gln Thr Glu Pro Glu Ser Arg Val Pro Val Leu Asp Phe Leu
 2245 2250 2255
 Glu Pro Met Ala Glu Glu Glu Ser Asp Leu Glu Pro Ser Ile Pro Ser
 2260 2265 2270
 Glu Cys Met Leu Pro Arg Ser Gly Phe Pro Arg Ala Leu Pro Ala Trp
 2275 2280 2285
 Ala Arg Pro Asp Tyr Asn Pro Pro Leu Val Glu Ser Trp Arg Arg Pro
 2290 2295 2300
 Asp Tyr Gln Pro Pro Thr Val Ala Gly Cys Ala Leu Pro Pro Pro Lys
 2305 2310 2315 2320
 Lys Ala Pro Thr Pro Pro Pro Arg Arg Arg Arg Thr Val Gly Leu Ser
 2325 2330 2335
 Glu Ser Thr Ile Ser Glu Ala Leu Gln Gln Leu Ala Ile Lys Thr Phe
 2340 2345 2350
 Gly Gln Pro Pro Ser Ser Gly Asp Ala Gly Ser Ser Thr Gly Ala Gly
 2355 2360 2365
 Ala Ala Glu Ser Gly Gly Pro Thr Ser Pro Gly Glu Pro Ala Pro Ser
 2370 2375 2380
 Glu Thr Gly Ser Ala Ser Ser Met Pro Pro Leu Glu Gly Glu Pro Gly
 2385 2390 2395 2400
 Asp Pro Asp Leu Glu Ser Asp Gln Val Glu Leu Gln Pro Pro Pro Gln
 2405 2410 2415
 Gly Gly Gly Val Ala Pro Gly Ser Gly Ser Gly Ser Trp Ser Thr Cys

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Ser Glu Glu Asp Asp Thr Thr Val Cys Cys Ser Met Ser Tyr Ser Trp		
2435	2440	2445
Thr Gly Ala Leu Ile Thr Pro Cys Ser Pro Glu Glu Glu Lys Leu Pro		
2450	2455	2460
Ile Asn Pro Leu Ser Asn Ser Leu Leu Arg Tyr His Asn Lys Val Tyr		
2465	2470	2475
Cys Thr Thr Ser Lys Ser Ala Ser Gln Arg Ala Lys Lys Val Thr Phe		
2485	2490	2495
Asp Arg Thr Gln Val Leu Asp Ala His Tyr Asp Ser Val Leu Lys Asp		
2500	2505	2510
Ile Lys Leu Ala Ala Ser Lys Val Ser Ala Arg Leu Leu Thr Leu Glu		
2515	2520	2525
Glu Ala Cys Gln Leu Thr Pro Pro His Ser Ala Arg Ser Lys Tyr Gly		
2530	2535	2540
Phe Gly Ala Lys Glu Val Arg Ser Leu Ser Gly Arg Ala Val Asn His		
2545	2550	2555
Ile Lys Ser Val Trp Lys Asp Leu Leu Glu Asp Pro Gln Thr Pro Ile		
2565	2570	2575
Pro Thr Thr Ile Met Ala Lys Asn Glu Val Phe Cys Val Asp Pro Ala		
2580	2585	2590
Lys Gly Gly Lys Lys Pro Ala Arg Leu Ile Val Tyr Pro Asp Leu Gly		
2595	2600	2605
Val Arg Val Cys Glu Lys Met Ala Leu Tyr Asp Ile Thr Gln Lys Leu		
2610	2615	2620
Pro Gln Ala Val Met Gly Ala Ser Tyr Gly Phe Gln Tyr Ser Pro Ala		
2625	2630	2635
Gln Arg Val Glu Tyr Leu Leu Lys Ala Trp Ala Glu Lys Lys Asp Pro		
2645	2650	2655

Met Gly Phe Ser Tyr Asp Thr Arg Cys Phe Asp Ser Thr Val Thr Glu
 2660 2665 2670
 Arg Asp Ile Arg Thr Glu Glu Ser Ile Tyr Gln Ala Cys Ser Leu Pro
 2675 2680 2685
 Glu Glu Ala Arg Thr Ala Ile His Ser Leu Thr Glu Arg Leu Tyr Val
 2690 2695 2700
 Gly Gly Pro Met Phe Asn Ser Lys Gly Gln Thr Cys Gly Tyr Arg Arg
 2705 2710 2715 2720
 Cys Arg Ala Ser Gly Val Leu Thr Thr Ser Met Gly Asn Thr Ile Thr
 2725 2730 2735
 Cys Tyr Val Lys Ala Leu Ala Ala Cys Lys Ala Ala Gly Ile Val Ala
 2740 2745 2750
 Pro Thr Met Leu Val Cys Gly Asp Asp Leu Val Val Ile Ser Glu Ser
 2755 2760 2765
 Gln Gly Thr Glu Glu Asp Glu Arg Asn Leu Arg Ala Phe Thr Glu Ala
 2770 2775 2780
 Met Thr Arg Tyr Ser Ala Pro Pro Gly Asp Pro Pro Arg Pro Glu Tyr
 2785 2790 2795 2800
 Asp Leu Glu Leu Ile Thr Ser Cys Ser Ser Asn Val Ser Val Ala Leu
 2805 2810 2815
 Gly Pro Arg Gly Arg Arg Arg Tyr Tyr Leu Thr Arg Asp Pro Thr Thr
 2820 2825 2830
 Pro Leu Ala Arg Ala Ala Trp Glu Thr Val Arg His Ser Pro Ile Asn
 2835 2840 2845
 Ser Trp Leu Gly Asn Ile Ile Gln Tyr Ala Pro Thr Ile Trp Val Arg
 2850 2855 2860
 Met Val Leu Met Thr His Phe Phe Ser Ile Leu Met Val Gln Asp Thr
 2865 2870 2875 2880
 Leu Asp Gln Asn Leu Asn Phe Glu Met Tyr Gly Ser Val Tyr Ser Val

2885	2890	2895
Asn Pro Leu Asp Leu Pro Ala Ile Ile Glu Arg Leu His Gly Leu Asp		
2900	2905	2910
Ala Phe Ser Met His Thr Tyr Ser His His Glu Leu Thr Arg Val Ala		
2915	2920	2925
Ser Ala Leu Arg Lys Leu Gly Ala Pro Pro Leu Arg Val Trp Lys Ser		
2930	2935	2940
Arg Ala Arg Ala Val Arg Ala Ser Leu Ile Ser Arg Gly Gly Lys Ala		
2945	2950	2955
Ala Val Cys Gly Arg Tyr Leu Phe Asn Trp Ala Val Lys Thr Lys Leu		
2965	2970	2975
Lys Leu Thr Pro Leu Pro Glu Ala Arg Leu Leu Asp Leu Ser Ser Trp		
2980	2985	2990
Phe Thr Val Gly Ala Gly Gly Gly Asp Ile Phe His Ser Val Ser Arg		
2995	3000	3005
Ala Arg Pro Arg Ser Leu Leu Phe Gly Leu Leu Leu Leu Phe Val Gly		
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Val Gly Leu Phe Leu Leu Pro Ala Arg		
3025	3030	

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<220>

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ccccctcccg ggagagccat agtgggtctgc ggaaccggtg agtacaccgg aattgccggg 180

aagactgggt cctttcttgg ataaaccac tctatgcccg gccatttggg cgtgcccccg 240

caagactgct agccgagtag cgttgggttg cgaaaggcct tgtggtactg cctgataggg 300

tgcttgcgag tgccccggga ggtctcgtag accgtgcacc atg agc aca aat ccc 355

Met Ser Thr Asn Pro

1 5

aaa cct caa aga aaa acc aaa aga aac act aac cgt cgc cca caa gac 403

Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn Arg Arg Pro Gln Asp

10 15 20

gtt aag ttt ccg ggc ggc ggc cag atc gtt ggc gga gta tac ttg ttg 451

Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly Gly Val Tyr Leu Leu

25 30 35

ccg cgc agg ggc ccc agg ttg ggt gtg cgc gcg aca agg aag gct tcg 499

Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala Thr Arg Lys Ala Ser

40 45 50

gag cgg tcc cag cca cgt ggg agg cgc cag ccc atc ccc aaa cat cgg 547

Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro Ile Pro Lys His Arg
55 60 65

cgc tcc act ggc aag tcc tgg ggg aag cca gga tac ccc tgg ccc ctg 595
Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly Tyr Pro Trp Pro Leu
70 75 80 85

tat ggg aat gag ggc ctc ggt tgg gca gga tgg ctc ctg tcc cct cga 643
Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp Leu Leu Ser Pro Arg
90 95 100

ggt tcc cgt ccc tca tgg ggc ccc aat gac ccc cgg cat agg tcg cgc 691
Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro Arg His Arg Ser Arg
105 110 115

aat gtg ggt aag gtc atc gat acc cta acg tgc ggc ttt gcc gac ctc 739
Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
120 125 130

ttg ggg tac gtc ccc gtc gta ggc gcc ccg ctt agt ggc gtt gcc agt 787
Leu Gly Tyr Val Pro Val Val Gly Ala Pro Leu Ser Gly Val Ala Ser
135 140 145

gct ctc gcg cac ggc gtg aga gtc ctg gag gac ggg gtt aat ttt gca 835
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Phe Ala
150 155 160 165

aca ggg aac tta cct ggt tgc tcc ttt tct atc ttc ttg ctg gcc cta 883
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu

170	175	180	
ctg tcc tgc atc act act ccg gtc tct gct gtc caa gtg aag aac acc			931
Leu Ser Cys Ile Thr Thr Pro Val Ser Ala Val Gln Val Lys Asn Thr			
185	190	195	
agc aac gcc tat atg gcg act aac gac tgt tcc aat gac agc atc act			979
Ser Asn Ala Tyr Met Ala Thr Asn Asp Cys Ser Asn Asp Ser Ile Thr			
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tgg cag ctt gag gcc gca gtc ctc cat gtc ccc ggg tgc gtc ccg tgc			1027
Trp Gln Leu Glu Ala Ala Val Leu His Val Pro Gly Cys Val Pro Cys			
215	220	225	
gag aaa atg ggg aac aca tca cgg tgc tgg ata cca gtc tca cca aac			1075
Glu Lys Met Gly Asn Thr Ser Arg Cys Trp Ile Pro Val Ser Pro Asn			
230	235	240	245
gtg gct gtg cgg cag cct ggc gcc ctc acg cgg ggc ttg cgg acg cac			1123
Val Ala Val Arg Gln Pro Gly Ala Leu Thr Arg Gly Leu Arg Thr His			
250	255	260	
atc gac atg gtc gtg ttg tcc gcc acg ctc tgc tcc gct ctc tac gtg			1171
Ile Asp Met Val Val Leu Ser Ala Thr Leu Cys Ser Ala Leu Tyr Val			
265	270	275	
ggg gac ctc tgt ggc ggg gtg atg ctc gcg tcc cag atg ttc att gtc			1219
Gly Asp Leu Cys Gly Gly Val Met Leu Ala Ser Gln Met Phe Ile Val			
280	285	290	

tcg ccg cag cac cac tgg ttc gtg cag gaa tgc aat tgc tcc atc tac 1267
 Ser Pro Gln His His Trp Phe Val Gln Glu Cys Asn Cys Ser Ile Tyr

295

300

305

cct ggc gcc atc act ggg cac cgt atg gca tgg gac atg atg atg aac 1315
 Pro Gly Ala Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn

310

315

320

325

tgg tcg ccc acg acc acc atg atc ctg gcg tac gtg atg cgc gtt ccc 1363
 Trp Ser Pro Thr Thr Thr Met Ile Leu Ala Tyr Val Met Arg Val Pro

330

335

340

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 Glu Val Ile Ile Asp Ile Ile Ser Gly Ala His Trp Gly Val Met Phe

345

350

355

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 Gly Leu Ala Tyr Phe Ser Met Gln Gly Ala Trp Ala Lys Val Val Val

360

365

370

atc ctc ctg ctg gcc tct ggg gtg gac gcg tac acc acc acg act ggg 1507
 Ile Leu Leu Leu Ala Ser Gly Val Asp Ala Tyr Thr Thr Thr Thr Gly

375

380

385

agc gct gct ggg cgc act acc agt agc ctg gcc agc gcc ttc tcc cct 1555
 Ser Ala Ala Gly Arg Thr Thr Ser Ser Leu Ala Ser Ala Phe Ser Pro

390

395

400

405

ggc gct cgg cag aac att cag ctc att aat acc aat ggt agc tgg cac 1603
 Gly Ala Arg Gln Asn Ile Gln Leu Ile Asn Thr Asn Gly Ser Trp His

410 415 420

atc aac cgc acc gcc ctg aat tgc aac gat tcc ttg cac acc ggc ttc 1651
 Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu His Thr Gly Phe

425 430 435

ttc acg gcc ctg ttc tac atc cat aag ttc aac tcg tcg gga tgt ccc 1699
 Phe Thr Ala Leu Phe Tyr Ile His Lys Phe Asn Ser Ser Gly Cys Pro

440 445 450

gag cgc ctg tcc gcc tgt cgc aac atc gag gac ttc cgg ata gga tgg 1747
 Glu Arg Leu Ser Ala Cys Arg Asn Ile Glu Asp Phe Arg Ile Gly Trp

455 460 465

ggc gcc ctg caa tac gac gac aat gtc acc aat cca gaa gat atg agg 1795
 Gly Ala Leu Gln Tyr Asp Asp Asn Val Thr Asn Pro Glu Asp Met Arg

470 475 480 485

cca tat tgc tgg cac tac cca cca aaa cag tgt ggc gta gtc ccc gca 1843
 Pro Tyr Cys Trp His Tyr Pro Pro Lys Gln Cys Gly Val Val Pro Ala

490 495 500

ggg acc gtg tgc ggc cca gtg tac tgt ttc acc cct agc ccg gtg gta 1891
 Gly Thr Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val Val

505 510 515

gtg ggc acg acc gat aga ctt gga gtg cct act tac acg tgg gga gag 1939

Val Gly Thr Thr Asp Arg Leu Gly Val Pro Thr Tyr Thr Trp Gly Glu	
520	525
530	
aat gag aca gat gtc ttc cta ttg aac agc acc cga cca ccg tcg ggg	1987
Asn Glu Thr Asp Val Phe Leu Leu Asn Ser Thr Arg Pro Pro Ser Gly	
535	540
545	
tca tgg ttt ggc tgc acg tgg atg aac tcc act ggc ttc acc aag acc	2035
Ser Trp Phe Gly Cys Thr Trp Met Asn Ser Thr Gly Phe Thr Lys Thr	
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560	565
tgc ggc gca cca ccc tgc cgc act aga gct gac ttc aat acc agc aca	2083
Cys Gly Ala Pro Pro Cys Arg Thr Arg Ala Asp Phe Asn Thr Ser Thr	
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580	
gat ctg ttg tgc ccc acg gac tgt ttt aga aaa cat cct gaa gcc act	2131
Asp Leu Leu Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr	
585	590
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Tyr Ile Lys Cys Gly Ser Gly Pro Trp Leu Thr Pro Lys Cys Leu Val	
600	605
610	
gac tac ccc tac agg ctc tgg cat tac cct tgc aca gtc aat tac tcc	2227
Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Tyr Ser	
615	620
625	
acc ttc aag atc agg atg tat gtg ggg gga gtt gag cac agg ctc atg	2275
Thr Phe Lys Ile Arg Met Tyr Val Gly Gly Val Glu His Arg Leu Met	

630	635	640	645	
gcc gcg tgc aat ttc act cgt ggg gat cgc tgc aac ttg gag gat agg				2323
Ala Ala Cys Asn Phe Thr Arg Gly Asp Arg Cys Asn Leu Glu Asp Arg				
	650	655	660	
gac aga agt caa cag act cct ctg ttg cac tcc acc acg gaa tgg gcc				2371
Asp Arg Ser Gln Gln Thr Pro Leu Leu His Ser Thr Thr Glu Trp Ala				
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att ttg ccc tgc tct ttc tca gac ttg ccc gct ttg tcg act ggt ctt				2419
Ile Leu Pro Cys Ser Phe Ser Asp Leu Pro Ala Leu Ser Thr Gly Leu				
	680	685	690	
ctc cac ctc cac caa aat atc gtg gac gta caa tat atg tat ggc ctg				2467
Leu His Leu His Gln Asn Ile Val Asp Val Gln Tyr Met Tyr Gly Leu				
	695	700	705	
tca cct gcc ctc aca caa tat atc gtt cga tgg gag tgg gta gta ctc				2515
Ser Pro Ala Leu Thr Gln Tyr Ile Val Arg Trp Glu Trp Val Val Leu				
710	715	720	725	
tta ttc ctg ctc cta gcg gac gcc agg gtc tgc gcc tgc ttg tgg atg				2563
Leu Phe Leu Leu Leu Ala Asp Ala Arg Val Cys Ala Cys Leu Trp Met				
	730	735	740	
ctc atc ttg ctg ggc caa gcc gaa gca gca ctg gag aag ctg gtc gtc				2611
Leu Ile Leu Leu Gly Gln Ala Glu Ala Ala Leu Glu Lys Leu Val Val				
	745	750	755	

ttg cac gct gcg agc gca gct agc tgc aat ggc ttc ctg tat ttt gtc	2659
Leu His Ala Ala Ser Ala Ala Ser Cys Asn Gly Phe Leu Tyr Phe Val	
760 765 770	
atc ttt ctc gtg gct gct tgg cac atc aag ggt agg gtg gtc ccc ttg	2707
Ile Phe Leu Val Ala Ala Trp His Ile Lys Gly Arg Val Val Pro Leu	
775 780 785	
gct gct tat tcc ctt act ggc ctg tgg ccg ttc tgc cta ctg ctc cta	2755
Ala Ala Tyr Ser Leu Thr Gly Leu Trp Pro Phe Cys Leu Leu Leu Leu	
790 795 800 805	
gca ctg ccc cag cag gct tac gcc tat gat gca tct gtg cac gga cag	2803
Ala Leu Pro Gln Gln Ala Tyr Ala Tyr Asp Ala Ser Val His Gly Gln	
810 815 820	
gtg ggc gcg gct ttg cta gta ctg att acc ctc ttt aca ctc acc ccg	2851
Val Gly Ala Ala Leu Leu Val Leu Ile Thr Leu Phe Thr Leu Thr Pro	
825 830 835	
ggg tat aag acc ctt ctc agc cag tcc ctg tgg tgg ttg tgc tat ctc	2899
Gly Tyr Lys Thr Leu Leu Ser Gln Ser Leu Trp Trp Leu Cys Tyr Leu	
840 845 850	
ctg acc ctg gcg gaa acc atg gtc cag gag tgg gca cca tcc atg cag	2947
Leu Thr Leu Ala Glu Thr Met Val Gln Glu Trp Ala Pro Ser Met Gln	
855 860 865	

gcg cgc ggc ggc cgt gat ggc atc ata tgg gcc gcc acc ata ttt tgc 2995
 Ala Arg Gly Gly Arg Asp Gly Ile Ile Trp Ala Ala Thr Ile Phe Cys
 870 875 880 885

ccg ggc gta gtg ttt gac ata acc aag tgg ctc tta gcg gtg ctt ggg 3043
 Pro Gly Val Val Phe Asp Ile Thr Lys Trp Leu Leu Ala Val Leu Gly
 890 895 900

cct ggt tac ctc cta aga ggt gct ttg acg cgc gtg cca tat ttc gtc 3091
 Pro Gly Tyr Leu Leu Arg Gly Ala Leu Thr Arg Val Pro Tyr Phe Val
 905 910 915

aga gcc cac gct ctg ctg aga atg tgc act atg gtg agg cac ctc gcg 3139
 Arg Ala His Ala Leu Leu Arg Met Cys Thr Met Val Arg His Leu Ala
 920 925 930

ggg ggt agg tac gtc cag atg gcg cta tta gcc ctt ggc agg tgg act 3187
 Gly Gly Arg Tyr Val Gln Met Ala Leu Leu Ala Leu Gly Arg Trp Thr
 935 940 945

ggc act tac atc tat gac cac ctc acc cct atg tcg gat tgg gct gct 3235
 Gly Thr Tyr Ile Tyr Asp His Leu Thr Pro Met Ser Asp Trp Ala Ala
 950 955 960 965

agc ggc ctg cgg gac ttg gcg gtc gct gtg gag cct atc atc ttc agt 3283
 Ser Gly Leu Arg Asp Leu Ala Val Ala Val Glu Pro Ile Ile Phe Ser
 970 975 980

ccg atg gag aag aaa gtc atc gtt tgg gga gcg gag acg gct gcg tgc 3331

Pro Met Glu Lys Lys Val Ile Val Trp Gly Ala Glu Thr Ala Ala Cys

985

990

995

ggg gac atc ttg cac gga ctt ccc gtg tcc gcc cga ctc ggt cgg gag 3379

Gly Asp Ile Leu His Gly Leu Pro Val Ser Ala Arg Leu Gly Arg Glu

1000

1005

1010

atc ctc ctt ggc cca gct gat ggc tac acc tcc aag ggg tgg aag ctt 3427

Ile Leu Leu Gly Pro Ala Asp Gly Tyr Thr Ser Lys Gly Trp Lys Leu

1015

1020

1025

ctc gcc ccc atc acc gct tac gcc cag cag aca cga ggt ctc ttg ggc 3475

Leu Ala Pro Ile Thr Ala Tyr Ala Gln Gln Thr Arg Gly Leu Leu Gly

1030

1035

1040

1045

tct ata gtg gtg agc atg acg ggg cgt gac aag aca gaa cag gcc ggg 3523

Ser Ile Val Val Ser Met Thr Gly Arg Asp Lys Thr Glu Gln Ala Gly

1050

1055

1060

gag gtc caa gtc ctg tcc aca gtc act cag tcc ttc ctc gga aca tcc 3571

Glu Val Gln Val Leu Ser Thr Val Thr Gln Ser Phe Leu Gly Thr Ser

1065

1070

1075

att tcg ggg gtc tta tgg act gtt tac cac gga gct ggc aac aag aca 3619

Ile Ser Gly Val Leu Trp Thr Val Tyr His Gly Ala Gly Asn Lys Thr

1080

1085

1090

cta gcc ggc tcg cgg ggc ccg gtc acg cag atg tac tcg agc gcc gag 3667

Leu Ala Gly Ser Arg Gly Pro Val Thr Gln Met Tyr Ser Ser Ala Glu

1095	1100	1105	
ggg gac ttg gtc ggg tgg ccc agc cct cct ggg acc aaa tct ttg gag			3715
Gly Asp Leu Val Gly Trp Pro Ser Pro Pro Gly Thr Lys Ser Leu Glu			
1110	1115	1120	1125
ccg tgt acg tgt gga gcg gtc gac ctg tat ttg gtc acg cgg aac gct			3763
Pro Cys Thr Cys Gly Ala Val Asp Leu Tyr Leu Val Thr Arg Asn Ala			
	1130	1135	1140
gat gtc atc ccg gct cga aga cgc ggg gac aag cgg gga gcg ctg ctc			3811
Asp Val Ile Pro Ala Arg Arg Arg Gly Asp Lys Arg Gly Ala Leu Leu			
	1145	1150	1155
tcc ccg aga ccc ctt tcg acc ttg aag ggg tcc tcg ggg gga cct gtg			3859
Ser Pro Arg Pro Leu Ser Thr Leu Lys Gly Ser Ser Gly Gly Pro Val			
	1160	1165	1170
ctt tgc cct agg ggc cac gct gtc gga atc ttc cgg gca gct gtg tgc			3907
Leu Cys Pro Arg Gly His Ala Val Gly Ile Phe Arg Ala Ala Val Cys			
1175	1180	1185	
tct cgg ggt gtg gct aag tcc ata gat ttc atc ccc gtt gag acg ctc			3955
Ser Arg Gly Val Ala Lys Ser Ile Asp Phe Ile Pro Val Glu Thr Leu			
1190	1195	1200	1205
gac atc gtc acg cgg tct ccc acc ttt agt gac aac agc aca cca cca			4003
Asp Ile Val Thr Arg Ser Pro Thr Phe Ser Asp Asn Ser Thr Pro Pro			
	1210	1215	1220

gct gtg ccc cag acc tat cag gtg ggg tac ttg cac gcc ccc act ggc 4051
 Ala Val Pro Gln Thr Tyr Gln Val Gly Tyr Leu His Ala Pro Thr Gly

1225

1230

1235

agt gga aaa agc acc aag gtc ccc gtc gcg tac gcc gcc cag ggg tat 4099
 Ser Gly Lys Ser Thr Lys Val Pro Val Ala Tyr Ala Ala Gln Gly Tyr

1240

1245

1250

aaa gtg ctg gtg ctc aat ccc tcg gtg gct gcc acc ctg gga ttt ggg 4147
 Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly

1255

1260

1265

gcg tac ttg tcc aag gca cat ggc atc aac ccc aac att agg act gga 4195
 Ala Tyr Leu Ser Lys Ala His Gly Ile Asn Pro Asn Ile Arg Thr Gly

1270

1275

1280

1285

gtc aga act gtg acg acc ggg gag ccc att aca tac tcc acg tat ggt 4243
 Val Arg Thr Val Thr Thr Gly Glu Pro Ile Thr Tyr Ser Thr Tyr Gly

1290

1295

1300

aaa ttc ctc gcc gat ggg ggc tgc gca ggc ggc gcc tat gac atc atc 4291
 Lys Phe Leu Ala Asp Gly Gly Cys Ala Gly Gly Ala Tyr Asp Ile Ile

1305

1310

1315

ata tgc gat gaa tgc cac tct gtg gat gct acc act att ctc ggc atc 4339
 Ile Cys Asp Glu Cys His Ser Val Asp Ala Thr Thr Ile Leu Gly Ile

1320

1325

1330

ggg aca gtc ctt gac caa gca gag aca gcc ggg gtc agg cta act gta 4387
 Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Val Arg Leu Thr Val
 1335 1340 1345

ctg gcc acg gcc acg ccc ccc ggg tcg gtg aca acc ccc cat ccc aat 4435
 Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn
 1350 1355 1360 1365

ata gag gag gta gcc ctc gga cag gag ggt gag atc ccc ttc tat ggg 4483
 Ile Glu Glu Val Ala Leu Gly Gln Glu Gly Glu Ile Pro Phe Tyr Gly
 1370 1375 1380

agg gcg ttt ccc ctg tct tac atc aag gga ggg agg cac ttg att ttc 4531
 Arg Ala Phe Pro Leu Ser Tyr Ile Lys Gly Gly Arg His Leu Ile Phe
 1385 1390 1395

tgc cac tca aag aaa aag tgt gac gag ctc gca acg gcc ctt cgg ggc 4579
 Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Thr Ala Leu Arg Gly
 1400 1405 1410

atg ggc ttg aac gct gtg gca tat tac aga ggg ttg gac gtc tcc ata 4627
 Met Gly Leu Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Ile
 1415 1420 1425

ata cca act caa gga gat gtg gtg gtc gtt gcc acc gac gcc ctc atg 4675
 Ile Pro Thr Gln Gly Asp Val Val Val Val Ala Thr Asp Ala Leu Met
 1430 1435 1440 1445

acg ggg tat act gga gac ttt gac tcc gtg atc gac tgc aac gta gcg 4723

Thr Gly Tyr Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Val Ala
1450 1455 1460

gtc acc cag gcc gta gac ttc agc ctg gac ccc acc ttc act ata acc 4771
Val Thr Gln Ala Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Thr
1465 1470 1475

aca cag act gtc ccg caa gac gct gtc tca cgt agt cag cgc cga ggg 4819
Thr Gln Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg Gly
1480 1485 1490

cgc acg ggt aga gga aga ctg ggc att tat agg tat gtt tcc act ggt 4867
Arg Thr Gly Arg Gly Arg Leu Gly Ile Tyr Arg Tyr Val Ser Thr Gly
1495 1500 1505

gag cga gcc tca gga atg ttt gac agt gta gta ctc tgt gag tgc tac 4915
Glu Arg Ala Ser Gly Met Phe Asp Ser Val Val Leu Cys Glu Cys Tyr
1510 1515 1520 1525

gac gca gga gct gct tgg tat gag ctc tca cca gtg gag acg acc gtc 4963
Asp Ala Gly Ala Ala Trp Tyr Glu Leu Ser Pro Val Glu Thr Thr Val
1530 1535 1540

agg ctc agg gcg tat ttc aac acg cct ggc ttg cct gtg tgc cag gac 5011
Arg Leu Arg Ala Tyr Phe Asn Thr Pro Gly Leu Pro Val Cys Gln Asp
1545 1550 1555

cac ctt gag ttt tgg gag gca gtt ttc acc ggc ctc aca cac ata gac 5059
His Leu Glu Phe Trp Glu Ala Val Phe Thr Gly Leu Thr His Ile Asp

1560	1565	1570	
gct cat ttc ctt tcc cag aca aag cag tcg ggg gaa aat ttc gca tac			5107
Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Phe Ala Tyr			
1575	1580	1585	
tta gta gcc tat cag gcc aca gtg tgc gcc agg gcc aaa gcg ccc ccc			5155
Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Lys Ala Pro Pro			
1590	1595	1600	1605
ccg tcc tgg gac gtc atg tgg aag tgc ttg act cga ctc aag ccc acg			5203
Pro Ser Trp Asp Val Met Trp Lys Cys Leu Thr Arg Leu Lys Pro Thr			
	1610	1615	1620
ctt gtg ggc cct aca cct ctc ctg tac cgt ttg ggc tct gtt acc aac			5251
Leu Val Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ser Val Thr Asn			
	1625	1630	1635
gag gtc acc ctt aca cac ccc gtg aca aaa tac atc gcc aca tgc atg			5299
Glu Val Thr Leu Thr His Pro Val Thr Lys Tyr Ile Ala Thr Cys Met			
	1640	1645	1650
caa gct gac ctc gag gtc atg acc agc acg tgg gtc ctg gct ggg gga			5347
Gln Ala Asp Leu Glu Val Met Thr Ser Thr Trp Val Leu Ala Gly Gly			
	1655	1660	1665
gtc tta gca gcc gtc gcc gcg tat tgc tta gcg acc ggg tgt gtt tcc			5395
Val Leu Ala Ala Val Ala Ala Tyr Cys Leu Ala Thr Gly Cys Val Ser			
1670	1675	1680	1685

atc att ggc cgt tta cac atc aac cag cga gct gtc gtc gct ccg gac 5443
 Ile Ile Gly Arg Leu His Ile Asn Gln Arg Ala Val Val Ala Pro Asp

1690 1695 1700

aag gag gtc ctc tat gag gct ttt gat gag atg gag gaa tgt gcc tcc 5491
 Lys Glu Val Leu Tyr Glu Ala Phe Asp Glu Met Glu Glu Cys Ala Ser

1705 1710 1715

aga gcg gct ctc ctt gaa gag ggg cag cgg ata gcc gag atg ctg aag 5539
 Arg Ala Ala Leu Leu Glu Glu Gly Gln Arg Ile Ala Glu Met Leu Lys

1720 1725 1730

tcc aag atc caa ggc tta ttg cag caa gcc tct aaa cag gcc cag gac 5587
 Ser Lys Ile Gln Gly Leu Leu Gln Gln Ala Ser Lys Gln Ala Gln Asp

1735 1740 1745

ata caa ccc gct gtg caa gct tcg tgg ccc aag atg gag caa ttc tgg 5635
 Ile Gln Pro Ala Val Gln Ala Ser Trp Pro Lys Met Glu Gln Phe Trp

1750 1755 1760 1765

gcc aaa cat atg tgg aac ttc ata agc ggc att cag tac ctc gca gga 5683
 Ala Lys His Met Trp Asn Phe Ile Ser Gly Ile Gln Tyr Leu Ala Gly

1770 1775 1780

ctg tca aca ctg cca ggg aac cct gct gtg gct tcc atg atg gca ttc 5731
 Leu Ser Thr Leu Pro Gly Asn Pro Ala Val Ala Ser Met Met Ala Phe

1785 1790 1795

agc gcc gcc ctc acc agt ccg ttg tca act agc acc acc atc ctt ctt 5779
 Ser Ala Ala Leu Thr Ser Pro Leu Ser Thr Ser Thr Thr Ile Leu Leu
 1800 1805 1810

aac att ctg ggg ggc tgg ctg gcg tcc caa att gcg cca ccc gcg ggg 5827
 Asn Ile Leu Gly Gly Trp Leu Ala Ser Gln Ile Ala Pro Pro Ala Gly
 1815 1820 1825

gcc act ggc ttt gtt gtc agt ggc ctg gtg gga gct gct gtt ggc agc 5875
 Ala Thr Gly Phe Val Val Ser Gly Leu Val Gly Ala Ala Val Gly Ser
 1830 1835 1840 1845

ata ggc ttg ggt aaa gtg ctg gtg gac atc ctg gca ggg tat ggt gcg 5923
 Ile Gly Leu Gly Lys Val Leu Val Asp Ile Leu Ala Gly Tyr Gly Ala
 1850 1855 1860

ggc att tcg ggg gcc ctc gtc gcg ttt aag atc atg tct ggc gag aag 5971
 Gly Ile Ser Gly Ala Leu Val Ala Phe Lys Ile Met Ser Gly Glu Lys
 1865 1870 1875

ccc tcc atg gag gat gtc atc aac ttg ctg cct ggg att ctg tct cca 6019
 Pro Ser Met Glu Asp Val Ile Asn Leu Leu Pro Gly Ile Leu Ser Pro
 1880 1885 1890

ggt gct ctg gtg gtg gga gtc atc tgc gcg gcc att ctg cgc cgc cat 6067
 Gly Ala Leu Val Val Gly Val Ile Cys Ala Ala Ile Leu Arg Arg His
 1895 1900 1905

gtg gga ccg ggg gaa ggc gcg gtc caa tgg atg aac agg ctt atc gcc 6115

Val Gly Pro Gly Glu Gly Ala Val Gln Trp Met Asn Arg Leu Ile Ala
1910 1915 1920 1925

ttc gct tcc aga gga aac cac gtc gcc cct act cac tac gtg acg gag 6163
Phe Ala Ser Arg Gly Asn His Val Ala Pro Thr His Tyr Val Thr Glu
1930 1935 1940

tgc gat gcg tcg cag cgt gtc acc caa ctg ctt ggc tct ctc act ata 6211
Ser Asp Ala Ser Gln Arg Val Thr Gln Leu Leu Gly Ser Leu Thr Ile
1945 1950 1955

act agt cta ctc agg aga ctt cac aac tgg atc act gag gat tgc ccc 6259
Thr Ser Leu Leu Arg Arg Leu His Asn Trp Ile Thr Glu Asp Cys Pro
1960 1965 1970

atc cca tgc gcc ggc tcg tgg ctc cgc gat gtg tgg gac tgg gtc tgt 6307
Ile Pro Cys Ala Gly Ser Trp Leu Arg Asp Val Trp Asp Trp Val Cys
1975 1980 1985

acc atc cta aca gac ttt aag aac tgg ctg acc tcc aag ctg ttc cca 6355
Thr Ile Leu Thr Asp Phe Lys Asn Trp Leu Thr Ser Lys Leu Phe Pro
1990 1995 2000 2005

aag atg cct ggc ctc ccc ttt atc tct tgc caa aag ggg tac aag ggc 6403
Lys Met Pro Gly Leu Pro Phe Ile Ser Cys Gln Lys Gly Tyr Lys Gly
2010 2015 2020

gtg tgg gcc ggc act ggc atc atg acc aca cga tgc ccc tgc ggc gcc 6451
Val Trp Ala Gly Thr Gly Ile Met Thr Thr Arg Cys Pro Cys Gly Ala

2025	2030	2035	
aac atc tct ggc aac gtc cgc ttg ggc tct atg aga atc aca gga ccc			6499
Asn Ile Ser Gly Asn Val Arg Leu Gly Ser Met Arg Ile Thr Gly Pro			
2040	2045	2050	
aaa acc tgc atg aac acc tgg cag ggg acc ttt cct atc aat tgt tat			6547
Lys Thr Cys Met Asn Thr Trp Gln Gly Thr Phe Pro Ile Asn Cys Tyr			
2055	2060	2065	
aca gaa ggc cag tgc ttg ccg aaa ccc gcg tta aac ttc aag acc gcc			6595
Thr Glu Gly Gln Cys Leu Pro Lys Pro Ala Leu Asn Phe Lys Thr Ala			
2070	2075	2080	2085
atc tgg aga gtg gcg gcc tca gag tac gcg gaa gtg acg cag cac gga			6643
Ile Trp Arg Val Ala Ala Ser Glu Tyr Ala Glu Val Thr Gln His Gly			
2090	2095	2100	
tca tat gcc tat ata aca ggg ctg acc act gac aac tta aaa gtc cct			6691
Ser Tyr Ala Tyr Ile Thr Gly Leu Thr Thr Asp Asn Leu Lys Val Pro			
2105	2110	2115	
tgc caa ctc ccc tct cca gag ttt ttc tct tgg gtg gac gga gta caa			6739
Cys Gln Leu Pro Ser Pro Glu Phe Phe Ser Trp Val Asp Gly Val Gln			
2120	2125	2130	
atc cat agg tcc gcc ccc aca cca aag ccg ttt ttc cgg gat gag gtc			6787
Ile His Arg Ser Ala Pro Thr Pro Lys Pro Phe Phe Arg Asp Glu Val			
2135	2140	2145	

tcg ttc agc gtt ggg ctc aat tca ttt gtc gtc ggg tct cag ctt ccc	6835
Ser Phe Ser Val Gly Leu Asn Ser Phe Val Val Gly Ser Gln Leu Pro	
2150 2155 2160 2165	
tgt gac cct gag ccc gac act gag gta gtg atg tcc atg cta aca gac	6883
Cys Asp Pro Glu Pro Asp Thr Glu Val Val Met Ser Met Leu Thr Asp	
2170 2175 2180	
cca tcc cat atc acg gcg gag gct gca gcg cgg cgt tta gcg cgg ggg	6931
Pro Ser His Ile Thr Ala Glu Ala Ala Ala Arg Arg Leu Ala Arg Gly	
2185 2190 2195	
tca ccc cca tct gag gca agc tcc tca gcg agc cag ctg tcg gcg cca	6979
Ser Pro Pro Ser Glu Ala Ser Ser Ser Ala Ser Gln Leu Ser Ala Pro	
2200 2205 2210	
tcg ctg cga gcc acc tgc acc acc cac ggt agg acc tat gat gtg gac	7027
Ser Leu Arg Ala Thr Cys Thr Thr His Gly Arg Thr Tyr Asp Val Asp	
2215 2220 2225	
atg gtg gat gcc aac ctg ttc atg ggg ggc ggc gtg att cgg ata gag	7075
Met Val Asp Ala Asn Leu Phe Met Gly Gly Gly Val Ile Arg Ile Glu	
2230 2235 2240 2245	
tct gag tcc aaa gtg gtc gtt ctg gac tcc ctc gac tca atg acc gag	7123
Ser Glu Ser Lys Val Val Val Leu Asp Ser Leu Asp Ser Met Thr Glu	
2250 2255 2260	

gaa gag ggc gac ctt gag cct tca gta cca tcg gag tat atg ctc ccc 7171
 Glu Glu Gly Asp Leu Glu Pro Ser Val Pro Ser Glu Tyr Met Leu Pro

2265

2270

2275

agg aag agg ttc cca ccg gcc tta ccg gct tgg gcg cgg cct gat tac 7219
 Arg Lys Arg Phe Pro Pro Ala Leu Pro Ala Trp Ala Arg Pro Asp Tyr

2280

2285

2290

aac cca ccg ctt gtg gaa tcg tgg aag agg cca gat tac caa cca ccc 7267
 Asn Pro Pro Leu Val Glu Ser Trp Lys Arg Pro Asp Tyr Gln Pro Pro

2295

2300

2305

act gtt gcg ggc tgt gct ctc ccc ccc ccc aaa aag acc ccg acg cct 7315
 Thr Val Ala Gly Cys Ala Leu Pro Pro Pro Lys Lys Thr Pro Thr Pro

2310

2315

2320

2325

cct cca agg aga cgc cgg aca gtg ggt ctg agc gag agc acc ata gga 7363
 Pro Pro Arg Arg Arg Arg Thr Val Gly Leu Ser Glu Ser Thr Ile Gly

2330

2335

2340

gat gcc ctc caa cag ctg gcc atc aag tcc ttt ggc cag ccc ccc cca 7411
 Asp Ala Leu Gln Gln Leu Ala Ile Lys Ser Phe Gly Gln Pro Pro Pro

2345

2350

2355

agc ggc gat tca ggc ctt tcc acg ggg gcg gac gcc gcc gac tcc ggc 7459
 Ser Gly Asp Ser Gly Leu Ser Thr Gly Ala Asp Ala Ala Asp Ser Gly

2360

2365

2370

gat cgg aca ccc cct gac gag ttg gct ctt tcg gag aca ggt tct acc 7507

Asp Arg Thr Pro Pro Asp Glu Leu Ala Leu Ser Glu Thr Gly Ser Thr
 2375 2380 2385

tcc tcc atg ccc ccc ctc gag ggg gag cct ggg gac cca gac ctg gag 7555
 Ser Ser Met Pro Pro Leu Glu Gly Glu Pro Gly Asp Pro Asp Leu Glu
 2390 2395 2400 2405

cct gag cag gta gag ctt caa cct cct ccc cag ggg ggg gag gca gct 7603
 Pro Glu Gln Val Glu Leu Gln Pro Pro Pro Gln Gly Gly Glu Ala Ala
 2410 2415 2420

ccc ggc tcg gac tcg ggg tcc tgg tct act tgc tcc gag gag gat gac 7651
 Pro Gly Ser Asp Ser Gly Ser Trp Ser Thr Cys Ser Glu Glu Asp Asp
 2425 2430 2435

tcc gtc gtg tgc tgc tcc atg tca tat tcc tgg acc ggg gct cta ata 7699
 Ser Val Val Cys Cys Ser Met Ser Tyr Ser Trp Thr Gly Ala Leu Ile
 2440 2445 2450

act cct tgt agc ccc gaa gag gaa aag ttg cca att aac tcc ttg agc 7747
 Thr Pro Cys Ser Pro Glu Glu Glu Lys Leu Pro Ile Asn Ser Leu Ser
 2455 2460 2465

aac tcg ctg ttg cga tac cat aac aag gta tac tgt act aca tca aag 7795
 Asn Ser Leu Leu Arg Tyr His Asn Lys Val Tyr Cys Thr Thr Ser Lys
 2470 2475 2480 2485

agt gcc tca cta agg gct aaa aag gta act ttt gat agg atg caa gtg 7843
 Ser Ala Ser Leu Arg Ala Lys Lys Val Thr Phe Asp Arg Met Gln Val

2490	2495	2500	
ctc gac gcc tat tat gat tca gtc tta aag gac atc aag cta gcg gcc			7891
Leu Asp Ala Tyr Tyr Asp Ser Val Leu Lys Asp Ile Lys Leu Ala Ala			
2505	2510	2515	
tcc aag gtc agc gca agg ctc ctc acc tta gag gag gcg tgc caa ttg			7939
Ser Lys Val Ser Ala Arg Leu Leu Thr Leu Glu Glu Ala Cys Gln Leu			
2520	2525	2530	
acc cca ccc cac tct gca aga tcc aag tat ggg ttt ggg gct aag gag			7987
Thr Pro Pro His Ser Ala Arg Ser Lys Tyr Gly Phe Gly Ala Lys Glu			
2535	2540	2545	
gtc cgc agc ttg tcc ggg agg gcc gtc aac cac atc aag tcc gtg tgg			8035
Val Arg Ser Leu Ser Gly Arg Ala Val Asn His Ile Lys Ser Val Trp			
2550	2555	2560	2565
aag gac ctc ttg gaa gac tca caa aca cca att cct aca acc atc atg			8083
Lys Asp Leu Leu Glu Asp Ser Gln Thr Pro Ile Pro Thr Thr Ile Met			
2570	2575	2580	
gcc aaa aat gag gtg ttc tgc gtg gac ccc gcc aag ggg ggt aaa aaa			8131
Ala Lys Asn Glu Val Phe Cys Val Asp Pro Ala Lys Gly Gly Lys Lys			
2585	2590	2595	
cca gct cgc ctt atc gtt tac cct gac ctc ggc gtc agg gtc tgc gag			8179
Pro Ala Arg Leu Ile Val Tyr Pro Asp Leu Gly Val Arg Val Cys Glu			
2600	2605	2610	

aag atg gcc ctt tat gat gtc aca caa aag ctt cct cag gcg gtg atg	8227
Lys Met Ala Leu Tyr Asp Val Thr Gln Lys Leu Pro Gln Ala Val Met	
2615	2620
2625	
ggg gct tct tat ggc ttc cag tac tcc ccc gct cag cgg gtg gag ttt	8275
Gly Ala Ser Tyr Gly Phe Gln Tyr Ser Pro Ala Gln Arg Val Glu Phe	
2630	2635
2640	2645
ctc ttg aag gca tgg gcg gaa aag aga gac cct atg ggt ttt tcg tat	8323
Leu Leu Lys Ala Trp Ala Glu Lys Arg Asp Pro Met Gly Phe Ser Tyr	
2650	2655
2660	
gat acc cga tgc ttt gac tca acc gtc act gag aga gac atc agg act	8371
Asp Thr Arg Cys Phe Asp Ser Thr Val Thr Glu Arg Asp Ile Arg Thr	
2665	2670
2675	
gag gag tcc ata tac cag gcc tgc tcc tta ccc gag gag gcc cga act	8419
Glu Glu Ser Ile Tyr Gln Ala Cys Ser Leu Pro Glu Glu Ala Arg Thr	
2680	2685
2690	
gcc ata cac tcg ctg act gag aga ctc tat gtg gga ggg ccc atg ttc	8467
Ala Ile His Ser Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Met Phe	
2695	2700
2705	
aac agc aag ggc cag tcc tgc ggg tac agg cgt tgc cgc gcc agc ggg	8515
Asn Ser Lys Gly Gln Ser Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly	
2710	2715
2720	2725

gtg ctt acc act agt atg ggg aac acc atc aca tgc tat gta aaa gcc 8563
Val Leu Thr Thr Ser Met Gly Asn Thr Ile Thr Cys Tyr Val Lys Ala

2730 2735 2740

cta gcg gct tgc aag gct gcg ggg ata att gcg ccc acg atg ctg gta 8611
Leu Ala Ala Cys Lys Ala Ala Gly Ile Ile Ala Pro Thr Met Leu Val

2745 2750 2755

tgc ggc gac gac ttg gtc gtc atc tca gaa agc cag ggg act gag gag 8659
Cys Gly Asp Asp Leu Val Val Ile Ser Glu Ser Gln Gly Thr Glu Glu

2760 2765 2770

gac gag cgg aac ctg aga gcc ttc acg gag gct atg acc agg tat tct 8707
Asp Glu Arg Asn Leu Arg Ala Phe Thr Glu Ala Met Thr Arg Tyr Ser

2775 2780 2785

gcc cct cct ggt gac ccc ccc aga ccg gaa tat gac ctg gag cta ata 8755
Ala Pro Pro Gly Asp Pro Pro Arg Pro Glu Tyr Asp Leu Glu Leu Ile

2790 2795 2800 2805

aca tct tgt tcc tca aac gtg tct gtg gca ctt ggc cca cag ggc cgc 8803
Thr Ser Cys Ser Ser Asn Val Ser Val Ala Leu Gly Pro Gln Gly Arg

2810 2815 2820

cgc aga tac tac ctg acc aga gac ccc acc act tca att gcc cgg gct 8851
Arg Arg Tyr Tyr Leu Thr Arg Asp Pro Thr Thr Ser Ile Ala Arg Ala

2825 2830 2835

gcc tgg gaa aca gtt aga cac tcc cct gtc aat tca tgg ctg gga aac 8899

Ala Trp Glu Thr Val Arg His Ser Pro Val Asn Ser Trp Leu Gly Asn	
2840	2845 2850
atc atc cag tac gct cca acc ata tgg gtt cgc atg gtc ctg atg aca	8947
Ile Ile Gln Tyr Ala Pro Thr Ile Trp Val Arg Met Val Leu Met Thr	
2855	2860 2865
cac ttc ttc tcc att ctc atg gcc cag gac acc cta gac cag aac ctt	8995
His Phe Phe Ser Ile Leu Met Ala Gln Asp Thr Leu Asp Gln Asn Leu	
2870	2875 2880 2885
aac ttt gaa atg tac gga tgc gtg tac tcc gtg agt cct ctg gac ctc	9043
Asn Phe Glu Met Tyr Gly Ser Val Tyr Ser Val Ser Pro Leu Asp Leu	
2890	2895 2900
cca gcc ata att gaa agg tta cac ggg ctt gac gcc ttc tct ctg cac	9091
Pro Ala Ile Ile Glu Arg Leu His Gly Leu Asp Ala Phe Ser Leu His	
2905	2910 2915
aca tac act ccc cac gaa ctg acg cgg gtg gct tca gcc ctc aga aaa	9139
Thr Tyr Thr Pro His Glu Leu Thr Arg Val Ala Ser Ala Leu Arg Lys	
2920	2925 2930
ctt ggg gcg cca ccc ctc aga gcg tgg aag agt cgg gcg cgt gca gtt	9187
Leu Gly Ala Pro Pro Leu Arg Ala Trp Lys Ser Arg Ala Arg Ala Val	
2935	2940 2945
agg gcg tcc ctc atc tcc cgt ggg ggg agg gcg gcc gtt tgc ggt cgg	9235
Arg Ala Ser Leu Ile Ser Arg Gly Gly Arg Ala Ala Val Cys Gly Arg	

2950	2955	2960	2965	
tac ctc ttc aac tgg gcg gtg aag acc aag ctc aaa ctc act cct ttg				9283
Tyr Leu Phe Asn Trp Ala Val Lys Thr Lys Leu Lys Leu Thr Pro Leu				
	2970	2975	2980	
ccg gag gca cgc ctc ctg gat ttg tcc agt tgg ttt acc gtc ggc gcc				9331
Pro Glu Ala Arg Leu Leu Asp Leu Ser Ser Trp Phe Thr Val Gly Ala				
	2985	2990	2995	
ggc ggg ggc gac att tat cac agc gtg tgc cgt gcc cga ccc cgc cta				9379
Gly Gly Gly Asp Ile Tyr His Ser Val Ser Arg Ala Arg Pro Arg Leu				
	3000	3005	3010	
tta ctc ctt agc cta ctc cta ctt tct gta ggg gta ggc ctc ttc cta				9427
Leu Leu Leu Ser Leu Leu Leu Leu Ser Val Gly Val Gly Leu Phe Leu				
	3015	3020	3025	
ctc ccc gct cga tag agcggcacac attagctaca ctccatagct aactgttcct				9482
Leu Pro Ala Arg				
3030				
tttttttttt tttttttttt tttttttttt ttttttttctt tttttttttt tttccctctt				9542
tcttcccttc tcatcttatt ctactttctt tcttggtggc tccatcttag ccctggtcac				9602
ggctagctgt gaaaggtccg tgagccgcat gactgcagag agtgccgtaa ctggtctctc				9662
tgcagatcat gt				9674

<210> 6

<211> 3033

<212> PRT

<213> Hepatitis C virus

<400> 6

Met	Ser	Thr	Asn	Pro	Lys	Pro	Gln	Arg	Lys	Thr	Lys	Arg	Asn	Thr	Asn
1				5					10					15	
Arg	Arg	Pro	Gln	Asp	Val	Lys	Phe	Pro	Gly	Gly	Gly	Gln	Ile	Val	Gly
			20					25					30		
Gly	Val	Tyr	Leu	Leu	Pro	Arg	Arg	Gly	Pro	Arg	Leu	Gly	Val	Arg	Ala
		35					40					45			
Thr	Arg	Lys	Ala	Ser	Glu	Arg	Ser	Gln	Pro	Arg	Gly	Arg	Arg	Gln	Pro
	50					55					60				
Ile	Pro	Lys	His	Arg	Arg	Ser	Thr	Gly	Lys	Ser	Trp	Gly	Lys	Pro	Gly
65				70						75				80	
Tyr	Pro	Trp	Pro	Leu	Tyr	Gly	Asn	Glu	Gly	Leu	Gly	Trp	Ala	Gly	Trp
			85						90					95	
Leu	Leu	Ser	Pro	Arg	Gly	Ser	Arg	Pro	Ser	Trp	Gly	Pro	Asn	Asp	Pro
		100						105					110		
Arg	His	Arg	Ser	Arg	Asn	Val	Gly	Lys	Val	Ile	Asp	Thr	Leu	Thr	Cys
		115					120					125			
Gly	Phe	Ala	Asp	Leu	Leu	Gly	Tyr	Val	Pro	Val	Val	Gly	Ala	Pro	Leu
	130					135						140			
Ser	Gly	Val	Ala	Ser	Ala	Leu	Ala	His	Gly	Val	Arg	Val	Leu	Glu	Asp
145					150					155				160	
Gly	Val	Asn	Phe	Ala	Thr	Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe	Ser	Ile

	165		170		175
Phe	Leu	Leu	Ala	Leu	Leu
		Ser	Cys	Ile	Thr
				Thr	Pro
				Val	Ser
				Ala	Val
	180		185		190
Gln	Val	Lys	Asn	Thr	Ser
		Asn	Ala	Tyr	Met
				Ala	Thr
				Asn	Asp
				Cys	Ser
	195		200		205
Asn	Asp	Ser	Ile	Thr	Trp
		Gln	Leu	Glu	Ala
				Ala	Val
				Leu	His
				Val	Pro
	210		215		220
Gly	Cys	Val	Pro	Cys	Glu
		Lys	Met	Gly	Asn
				Thr	Ser
				Arg	Cys
				Trp	Ile
	225		230		235
					240
Pro	Val	Ser	Pro	Asn	Val
				Ala	Val
				Arg	Gln
				Pro	Gly
				Ala	Leu
				Thr	Arg
	245		250		255
Gly	Leu	Arg	Thr	His	Ile
		Asp	Met	Val	Val
				Leu	Ser
				Ala	Thr
				Leu	Cys
	260		265		270
Ser	Ala	Leu	Tyr	Val	Gly
		Asp	Leu	Cys	Gly
				Gly	Val
				Met	Leu
				Ala	Ser
	275		280		285
Gln	Met	Phe	Ile	Val	Ser
				Pro	Gln
				His	His
				Trp	Phe
				Val	Gln
				Glu	Cys
	290		295		300
Asn	Cys	Ser	Ile	Tyr	Pro
				Gly	Ala
				Ile	Thr
				Gly	His
				Arg	Met
				Ala	Trp
	305		310		315
					320
Asp	Met	Met	Met	Asn	Trp
				Ser	Pro
				Thr	Thr
				Thr	Met
				Ile	Leu
				Ala	Tyr
	325		330		335
Val	Met	Arg	Val	Pro	Glu
				Val	Ile
				Ile	Asp
				Ile	Ile
				Ser	Gly
				Ala	His
	340		345		350
Trp	Gly	Val	Met	Phe	Gly
				Leu	Ala
				Tyr	Phe
				Ser	Met
				Gln	Gly
				Ala	Trp
	355		360		365
Ala	Lys	Val	Val	Val	Ile
				Leu	Leu
				Leu	Ala
				Ser	Gly
				Val	Asp
				Ala	Tyr
	370		375		380
Thr	Thr	Thr	Thr	Gly	Ser
				Ala	Ala
				Gly	Arg
				Thr	Thr
				Ser	Ser
				Leu	Ala
	385		390		395
					400

Ser Ala Phe Ser Pro Gly Ala Arg Gln Asn Ile Gln Leu Ile Asn Thr
 405 410 415
 Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser
 420 425 430
 Leu His Thr Gly Phe Phe Thr Ala Leu Phe Tyr Ile His Lys Phe Asn
 435 440 445
 Ser Ser Gly Cys Pro Glu Arg Leu Ser Ala Cys Arg Asn Ile Glu Asp
 450 455 460
 Phe Arg Ile Gly Trp Gly Ala Leu Gln Tyr Asp Asp Asn Val Thr Asn
 465 470 475 480
 Pro Glu Asp Met Arg Pro Tyr Cys Trp His Tyr Pro Pro Lys Gln Cys
 485 490 495
 Gly Val Val Pro Ala Gly Thr Val Cys Gly Pro Val Tyr Cys Phe Thr
 500 505 510
 Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Leu Gly Val Pro Thr
 515 520 525
 Tyr Thr Trp Gly Glu Asn Glu Thr Asp Val Phe Leu Leu Asn Ser Thr
 530 535 540
 Arg Pro Pro Ser Gly Ser Trp Phe Gly Cys Thr Trp Met Asn Ser Thr
 545 550 555 560
 Gly Phe Thr Lys Thr Cys Gly Ala Pro Pro Cys Arg Thr Arg Ala Asp
 565 570 575
 Phe Asn Thr Ser Thr Asp Leu Leu Cys Pro Thr Asp Cys Phe Arg Lys
 580 585 590
 His Pro Glu Ala Thr Tyr Ile Lys Cys Gly Ser Gly Pro Trp Leu Thr
 595 600 605
 Pro Lys Cys Leu Val Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys
 610 615 620
 Thr Val Asn Tyr Ser Thr Phe Lys Ile Arg Met Tyr Val Gly Gly Val

625	630	635	640
Glu His Arg Leu Met Ala Ala Cys Asn Phe Thr Arg Gly Asp Arg Cys			
645	650	655	
Asn Leu Glu Asp Arg Asp Arg Ser Gln Gln Thr Pro Leu Leu His Ser			
660	665	670	
Thr Thr Glu Trp Ala Ile Leu Pro Cys Ser Phe Ser Asp Leu Pro Ala			
675	680	685	
Leu Ser Thr Gly Leu Leu His Leu His Gln Asn Ile Val Asp Val Gln			
690	695	700	
Tyr Met Tyr Gly Leu Ser Pro Ala Leu Thr Gln Tyr Ile Val Arg Trp			
705	710	715	720
Glu Trp Val Val Leu Leu Phe Leu Leu Leu Ala Asp Ala Arg Val Cys			
725	730	735	
Ala Cys Leu Trp Met Leu Ile Leu Leu Gly Gln Ala Glu Ala Ala Leu			
740	745	750	
Glu Lys Leu Val Val Leu His Ala Ala Ser Ala Ala Ser Cys Asn Gly			
755	760	765	
Phe Leu Tyr Phe Val Ile Phe Leu Val Ala Ala Trp His Ile Lys Gly			
770	775	780	
Arg Val Val Pro Leu Ala Ala Tyr Ser Leu Thr Gly Leu Trp Pro Phe			
785	790	795	800
Cys Leu Leu Leu Leu Ala Leu Pro Gln Gln Ala Tyr Ala Tyr Asp Ala			
805	810	815	
Ser Val His Gly Gln Val Gly Ala Ala Leu Leu Val Leu Ile Thr Leu			
820	825	830	
Phe Thr Leu Thr Pro Gly Tyr Lys Thr Leu Leu Ser Gln Ser Leu Trp			
835	840	845	
Trp Leu Cys Tyr Leu Leu Thr Leu Ala Glu Thr Met Val Gln Glu Trp			
850	855	860	

Ala Pro Ser Met Gln Ala Arg Gly Gly Arg Asp Gly Ile Ile Trp Ala			
865	870	875	880
Ala Thr Ile Phe Cys Pro Gly Val Val Phe Asp Ile Thr Lys Trp Leu			
	885	890	895
Leu Ala Val Leu Gly Pro Gly Tyr Leu Leu Arg Gly Ala Leu Thr Arg			
	900	905	910
Val Pro Tyr Phe Val Arg Ala His Ala Leu Leu Arg Met Cys Thr Met			
	915	920	925
Val Arg His Leu Ala Gly Gly Arg Tyr Val Gln Met Ala Leu Leu Ala			
	930	935	940
Leu Gly Arg Trp Thr Gly Thr Tyr Ile Tyr Asp His Leu Thr Pro Met			
945	950	955	960
Ser Asp Trp Ala Ala Ser Gly Leu Arg Asp Leu Ala Val Ala Val Glu			
	965	970	975
Pro Ile Ile Phe Ser Pro Met Glu Lys Lys Val Ile Val Trp Gly Ala			
	980	985	990
Glu Thr Ala Ala Cys Gly Asp Ile Leu His Gly Leu Pro Val Ser Ala			
	995	1000	1005
Arg Leu Gly Arg Glu Ile Leu Leu Gly Pro Ala Asp Gly Tyr Thr Ser			
	1010	1015	1020
Lys Gly Trp Lys Leu Leu Ala Pro Ile Thr Ala Tyr Ala Gln Gln Thr			
1025	1030	1035	1040
Arg Gly Leu Leu Gly Ser Ile Val Val Ser Met Thr Gly Arg Asp Lys			
	1045	1050	1055
Thr Glu Gln Ala Gly Glu Val Gln Val Leu Ser Thr Val Thr Gln Ser			
	1060	1065	1070
Phe Leu Gly Thr Ser Ile Ser Gly Val Leu Trp Thr Val Tyr His Gly			
	1075	1080	1085
Ala Gly Asn Lys Thr Leu Ala Gly Ser Arg Gly Pro Val Thr Gln Met			

1090	1095	1100
Tyr Ser Ser Ala Glu Gly Asp Leu Val Gly Trp Pro Ser Pro Pro Gly		
1105	1110	1115
Thr Lys Ser Leu Glu Pro Cys Thr Cys Gly Ala Val Asp Leu Tyr Leu		
1125	1130	1135
Val Thr Arg Asn Ala Asp Val Ile Pro Ala Arg Arg Arg Gly Asp Lys		
1140	1145	1150
Arg Gly Ala Leu Leu Ser Pro Arg Pro Leu Ser Thr Leu Lys Gly Ser		
1155	1160	1165
Ser Gly Gly Pro Val Leu Cys Pro Arg Gly His Ala Val Gly Ile Phe		
1170	1175	1180
Arg Ala Ala Val Cys Ser Arg Gly Val Ala Lys Ser Ile Asp Phe Ile		
1185	1190	1195
Pro Val Glu Thr Leu Asp Ile Val Thr Arg Ser Pro Thr Phe Ser Asp		
1205	1210	1215
Asn Ser Thr Pro Pro Ala Val Pro Gln Thr Tyr Gln Val Gly Tyr Leu		
1220	1225	1230
His Ala Pro Thr Gly Ser Gly Lys Ser Thr Lys Val Pro Val Ala Tyr		
1235	1240	1245
Ala Ala Gln Gly Tyr Lys Val Leu Val Leu Asn Pro Ser Val Ala Ala		
1250	1255	1260
Thr Leu Gly Phe Gly Ala Tyr Leu Ser Lys Ala His Gly Ile Asn Pro		
1265	1270	1275
Asn Ile Arg Thr Gly Val Arg Thr Val Thr Thr Gly Glu Pro Ile Thr		
1285	1290	1295
Tyr Ser Thr Tyr Gly Lys Phe Leu Ala Asp Gly Gly Cys Ala Gly Gly		
1300	1305	1310
Ala Tyr Asp Ile Ile Ile Cys Asp Glu Cys His Ser Val Asp Ala Thr		
1315	1320	1325

Thr Ile Leu Gly Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly
 1330 1335 1340
 Val Arg Leu Thr Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr
 1345 1350 1355 1360
 Thr Pro His Pro Asn Ile Glu Glu Val Ala Leu Gly Gln Glu Gly Glu
 1365 1370 1375
 Ile Pro Phe Tyr Gly Arg Ala Phe Pro Leu Ser Tyr Ile Lys Gly Gly
 1380 1385 1390
 Arg His Leu Ile Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala
 1395 1400 1405
 Thr Ala Leu Arg Gly Met Gly Leu Asn Ala Val Ala Tyr Tyr Arg Gly
 1410 1415 1420
 Leu Asp Val Ser Ile Ile Pro Thr Gln Gly Asp Val Val Val Val Ala
 1425 1430 1435 1440
 Thr Asp Ala Leu Met Thr Gly Tyr Thr Gly Asp Phe Asp Ser Val Ile
 1445 1450 1455
 Asp Cys Asn Val Ala Val Thr Gln Ala Val Asp Phe Ser Leu Asp Pro
 1460 1465 1470
 Thr Phe Thr Ile Thr Thr Gln Thr Val Pro Gln Asp Ala Val Ser Arg
 1475 1480 1485
 Ser Gln Arg Arg Gly Arg Thr Gly Arg Gly Arg Leu Gly Ile Tyr Arg
 1490 1495 1500
 Tyr Val Ser Thr Gly Glu Arg Ala Ser Gly Met Phe Asp Ser Val Val
 1505 1510 1515 1520
 Leu Cys Glu Cys Tyr Asp Ala Gly Ala Ala Trp Tyr Glu Leu Ser Pro
 1525 1530 1535
 Val Glu Thr Thr Val Arg Leu Arg Ala Tyr Phe Asn Thr Pro Gly Leu
 1540 1545 1550
 Pro Val Cys Gln Asp His Leu Glu Phe Trp Glu Ala Val Phe Thr Gly

1555	1560	1565
Leu Thr His Ile Asp Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly		
1570	1575	1580
Glu Asn Phe Ala Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg		
1585	1590	1595
Ala Lys Ala Pro Pro Pro Ser Trp Asp Val Met Trp Lys Cys Leu Thr		1600
1605	1610	1615
Arg Leu Lys Pro Thr Leu Val Gly Pro Thr Pro Leu Leu Tyr Arg Leu		
1620	1625	1630
Gly Ser Val Thr Asn Glu Val Thr Leu Thr His Pro Val Thr Lys Tyr		
1635	1640	1645
Ile Ala Thr Cys Met Gln Ala Asp Leu Glu Val Met Thr Ser Thr Trp		
1650	1655	1660
Val Leu Ala Gly Gly Val Leu Ala Ala Val Ala Ala Tyr Cys Leu Ala		
1665	1670	1675
Thr Gly Cys Val Ser Ile Ile Gly Arg Leu His Ile Asn Gln Arg Ala		1680
1685	1690	1695
Val Val Ala Pro Asp Lys Glu Val Leu Tyr Glu Ala Phe Asp Glu Met		
1700	1705	1710
Glu Glu Cys Ala Ser Arg Ala Ala Leu Leu Glu Glu Gly Gln Arg Ile		
1715	1720	1725
Ala Glu Met Leu Lys Ser Lys Ile Gln Gly Leu Leu Gln Gln Ala Ser		
1730	1735	1740
Lys Gln Ala Gln Asp Ile Gln Pro Ala Val Gln Ala Ser Trp Pro Lys		
1745	1750	1755
Met Glu Gln Phe Trp Ala Lys His Met Trp Asn Phe Ile Ser Gly Ile		1760
1765	1770	1775
Gln Tyr Leu Ala Gly Leu Ser Thr Leu Pro Gly Asn Pro Ala Val Ala		
1780	1785	1790

Ser Met Met Ala Phe Ser Ala Ala Leu Thr Ser Pro Leu Ser Thr Ser
 1795 1800 1805
 Thr Thr Ile Leu Leu Asn Ile Leu Gly Gly Trp Leu Ala Ser Gln Ile
 1810 1815 1820
 Ala Pro Pro Ala Gly Ala Thr Gly Phe Val Val Ser Gly Leu Val Gly
 1825 1830 1835 1840
 Ala Ala Val Gly Ser Ile Gly Leu Gly Lys Val Leu Val Asp Ile Leu
 1845 1850 1855
 Ala Gly Tyr Gly Ala Gly Ile Ser Gly Ala Leu Val Ala Phe Lys Ile
 1860 1865 1870
 Met Ser Gly Glu Lys Pro Ser Met Glu Asp Val Ile Asn Leu Leu Pro
 1875 1880 1885
 Gly Ile Leu Ser Pro Gly Ala Leu Val Val Gly Val Ile Cys Ala Ala
 1890 1895 1900
 Ile Leu Arg Arg His Val Gly Pro Gly Glu Gly Ala Val Gln Trp Met
 1905 1910 1915 1920
 Asn Arg Leu Ile Ala Phe Ala Ser Arg Gly Asn His Val Ala Pro Thr
 1925 1930 1935
 His Tyr Val Thr Glu Ser Asp Ala Ser Gln Arg Val Thr Gln Leu Leu
 1940 1945 1950
 Gly Ser Leu Thr Ile Thr Ser Leu Leu Arg Arg Leu His Asn Trp Ile
 1955 1960 1965
 Thr Glu Asp Cys Pro Ile Pro Cys Ala Gly Ser Trp Leu Arg Asp Val
 1970 1975 1980
 Trp Asp Trp Val Cys Thr Ile Leu Thr Asp Phe Lys Asn Trp Leu Thr
 1985 1990 1995 2000
 Ser Lys Leu Phe Pro Lys Met Pro Gly Leu Pro Phe Ile Ser Cys Gln
 2005 2010 2015
 Lys Gly Tyr Lys Gly Val Trp Ala Gly Thr Gly Ile Met Thr Thr Arg

2020	2025	2030
Cys Pro Cys Gly Ala Asn Ile Ser Gly Asn Val Arg Leu Gly Ser Met		
2035	2040	2045
Arg Ile Thr Gly Pro Lys Thr Cys Met Asn Thr Trp Gln Gly Thr Phe		
2050	2055	2060
Pro Ile Asn Cys Tyr Thr Glu Gly Gln Cys Leu Pro Lys Pro Ala Leu		
2065	2070	2075
Asn Phe Lys Thr Ala Ile Trp Arg Val Ala Ala Ser Glu Tyr Ala Glu		
2085	2090	2095
Val Thr Gln His Gly Ser Tyr Ala Tyr Ile Thr Gly Leu Thr Thr Asp		
2100	2105	2110
Asn Leu Lys Val Pro Cys Gln Leu Pro Ser Pro Glu Phe Phe Ser Trp		
2115	2120	2125
Val Asp Gly Val Gln Ile His Arg Ser Ala Pro Thr Pro Lys Pro Phe		
2130	2135	2140
Phe Arg Asp Glu Val Ser Phe Ser Val Gly Leu Asn Ser Phe Val Val		
2145	2150	2155
Gly Ser Gln Leu Pro Cys Asp Pro Glu Pro Asp Thr Glu Val Val Met		
2165	2170	2175
Ser Met Leu Thr Asp Pro Ser His Ile Thr Ala Glu Ala Ala Ala Arg		
2180	2185	2190
Arg Leu Ala Arg Gly Ser Pro Pro Ser Glu Ala Ser Ser Ser Ala Ser		
2195	2200	2205
Gln Leu Ser Ala Pro Ser Leu Arg Ala Thr Cys Thr Thr His Gly Arg		
2210	2215	2220
Thr Tyr Asp Val Asp Met Val Asp Ala Asn Leu Phe Met Gly Gly Gly		
2225	2230	2235
Val Ile Arg Ile Glu Ser Glu Ser Lys Val Val Val Leu Asp Ser Leu		
2245	2250	2255

Asp Ser Met Thr Glu Glu Glu Gly Asp Leu Glu Pro Ser Val Pro Ser
 2260 2265 2270
 Glu Tyr Met Leu Pro Arg Lys Arg Phe Pro Pro Ala Leu Pro Ala Trp
 2275 2280 2285
 Ala Arg Pro Asp Tyr Asn Pro Pro Leu Val Glu Ser Trp Lys Arg Pro
 2290 2295 2300
 Asp Tyr Gln Pro Pro Thr Val Ala Gly Cys Ala Leu Pro Pro Pro Lys
 2305 2310 2315 2320
 Lys Thr Pro Thr Pro Pro Pro Arg Arg Arg Arg Thr Val Gly Leu Ser
 2325 2330 2335
 Glu Ser Thr Ile Gly Asp Ala Leu Gln Gln Leu Ala Ile Lys Ser Phe
 2340 2345 2350
 Gly Gln Pro Pro Pro Ser Gly Asp Ser Gly Leu Ser Thr Gly Ala Asp
 2355 2360 2365
 Ala Ala Asp Ser Gly Asp Arg Thr Pro Pro Asp Glu Leu Ala Leu Ser
 2370 2375 2380
 Glu Thr Gly Ser Thr Ser Ser Met Pro Pro Leu Glu Gly Glu Pro Gly
 2385 2390 2395 2400
 Asp Pro Asp Leu Glu Pro Glu Gln Val Glu Leu Gln Pro Pro Pro Gln
 2405 2410 2415
 Gly Gly Glu Ala Ala Pro Gly Ser Asp Ser Gly Ser Trp Ser Thr Cys
 2420 2425 2430
 Ser Glu Glu Asp Asp Ser Val Val Cys Cys Ser Met Ser Tyr Ser Trp
 2435 2440 2445
 Thr Gly Ala Leu Ile Thr Pro Cys Ser Pro Glu Glu Glu Lys Leu Pro
 2450 2455 2460
 Ile Asn Ser Leu Ser Asn Ser Leu Leu Arg Tyr His Asn Lys Val Tyr
 2465 2470 2475 2480
 Cys Thr Thr Ser Lys Ser Ala Ser Leu Arg Ala Lys Lys Val Thr Phe

2485	2490	2495
Asp Arg Met Gln Val Leu Asp Ala Tyr Tyr Asp Ser Val Leu Lys Asp		
2500	2505	2510
Ile Lys Leu Ala Ala Ser Lys Val Ser Ala Arg Leu Leu Thr Leu Glu		
2515	2520	2525
Glu Ala Cys Gln Leu Thr Pro Pro His Ser Ala Arg Ser Lys Tyr Gly		
2530	2535	2540
Phe Gly Ala Lys Glu Val Arg Ser Leu Ser Gly Arg Ala Val Asn His		
2545	2550	2555
Ile Lys Ser Val Trp Lys Asp Leu Leu Glu Asp Ser Gln Thr Pro Ile		
2565	2570	2575
Pro Thr Thr Ile Met Ala Lys Asn Glu Val Phe Cys Val Asp Pro Ala		
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Lys Gly Gly Lys Lys Pro Ala Arg Leu Ile Val Tyr Pro Asp Leu Gly		
2595	2600	2605
Val Arg Val Cys Glu Lys Met Ala Leu Tyr Asp Val Thr Gln Lys Leu		
2610	2615	2620
Pro Gln Ala Val Met Gly Ala Ser Tyr Gly Phe Gln Tyr Ser Pro Ala		
2625	2630	2635
Gln Arg Val Glu Phe Leu Leu Lys Ala Trp Ala Glu Lys Arg Asp Pro		
2645	2650	2655
Met Gly Phe Ser Tyr Asp Thr Arg Cys Phe Asp Ser Thr Val Thr Glu		
2660	2665	2670
Arg Asp Ile Arg Thr Glu Glu Ser Ile Tyr Gln Ala Cys Ser Leu Pro		
2675	2680	2685
Glu Glu Ala Arg Thr Ala Ile His Ser Leu Thr Glu Arg Leu Tyr Val		
2690	2695	2700
Gly Gly Pro Met Phe Asn Ser Lys Gly Gln Ser Cys Gly Tyr Arg Arg		
2705	2710	2715
		2720

Cys Arg Ala Ser Gly Val Leu Thr Thr Ser Met Gly Asn Thr Ile Thr
2725 2730 2735
Cys Tyr Val Lys Ala Leu Ala Ala Cys Lys Ala Ala Gly Ile Ile Ala
2740 2745 2750
Pro Thr Met Leu Val Cys Gly Asp Asp Leu Val Val Ile Ser Glu Ser
2755 2760 2765
Gln Gly Thr Glu Glu Asp Glu Arg Asn Leu Arg Ala Phe Thr Glu Ala
2770 2775 2780
Met Thr Arg Tyr Ser Ala Pro Pro Gly Asp Pro Pro Arg Pro Glu Tyr
2785 2790 2795 2800
Asp Leu Glu Leu Ile Thr Ser Cys Ser Ser Asn Val Ser Val Ala Leu
2805 2810 2815
Gly Pro Gln Gly Arg Arg Arg Tyr Tyr Leu Thr Arg Asp Pro Thr Thr
2820 2825 2830
Ser Ile Ala Arg Ala Ala Trp Glu Thr Val Arg His Ser Pro Val Asn
2835 2840 2845
Ser Trp Leu Gly Asn Ile Ile Gln Tyr Ala Pro Thr Ile Trp Val Arg
2850 2855 2860
Met Val Leu Met Thr His Phe Phe Ser Ile Leu Met Ala Gln Asp Thr
2865 2870 2875 2880
Leu Asp Gln Asn Leu Asn Phe Glu Met Tyr Gly Ser Val Tyr Ser Val
2885 2890 2895
Ser Pro Leu Asp Leu Pro Ala Ile Ile Glu Arg Leu His Gly Leu Asp
2900 2905 2910
Ala Phe Ser Leu His Thr Tyr Thr Pro His Glu Leu Thr Arg Val Ala
2915 2920 2925
Ser Ala Leu Arg Lys Leu Gly Ala Pro Pro Leu Arg Ala Trp Lys Ser
2930 2935 2940
Arg Ala Arg Ala Val Arg Ala Ser Leu Ile Ser Arg Gly Gly Arg Ala

2945	2950	2955	2960
Ala Val Cys Gly Arg Tyr Leu Phe Asn Trp	Ala Val Lys Thr Lys Leu		
2965	2970	2975	
Lys Leu Thr Pro Leu Pro Glu Ala Arg Leu Leu Asp Leu Ser Ser Trp			
2980	2985	2990	
Phe Thr Val Gly Ala Gly Gly Gly Asp Ile Tyr His Ser Val Ser Arg			
2995	3000	3005	
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<211> 340

<212> RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic RNA

<400> 9

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 ccccccuccg ggagagccau aguggucugc ggaaccggug aguacaccgg aaugccggg 180
 aagacugggu ccuuucuuu auaaaccac ucuaucccg gccauuuggg cgugccccc 240
 caagacugcu agccgaguag cguuggguug cgaaaggccu ugugguacug ccugauaggg 300
 cguugcgag ugccccggga ggucucguag accugcacc 340

<210> 10

<211> 340

<212> RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic RNA

<400> 10

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cuucacgcag aaagcgucua gccauggcgu uaguaugagu gucguacagc cuccaggccc 120
ccccucccg ggagagccau aguggucugc ggaaccggug aguacaccgg aauugccggg 180
aagacugggu ccuuucuugg aaaaaccac ucuaugcccg gccauuuggg cgugcccccg 240
caagacugcu agccgaguag cguuggguug cgaaaggccu ugugguacug ccugauaggg 300
ugcuugcgag ugccccggga ggucucguag accgugcacc 340
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<210> 11

<211> 236

<212> RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic RNA

<400> 11

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agcggcacac acuagguaca cuccauagcu aacuguuccu uuuuuuuuuu uuuuuuuuuu 60
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```

uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu 120
uauuucuacuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu 180
uccgugagcc gcaugacugc agagagugcc gaaacugguc ucucugcaga ucaugu 236

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<210> 12

<211> 232

<212> RNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic RNA

<400> 12

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agcggcacac auuagcuaca cuccauagcu aacuguuccu uuuuuuuuuuuu uuuuuuuuuuuu 60
uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu uuuuuuuuuuuu 120
cuacuuuucuu ucuugguggc uccaucuuag ccugugucac ggcuaugcugu gaaagguccg 180
ugagccgcgau gacugcagag agugccguua cuggucucuc ugcagaucuu gu 232

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<210> 13

<211> 17

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 13

cgggagagcc atagtgg

17

<210> 14

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 14

agtaccacaa ggcctttcg

19

<210> 15

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 15

ctgcggaacc ggtgagtaca c

21

<210> 16

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 16

aacaagatgg attgcacgca

20

<210> 17

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 17

cgtcaagaag gcgatagaag

20

<210> 18

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 18
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<210> 19

<211> 28

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic DNA

<400> 19
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<210> 20

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 20
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<210> 21

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 21

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30

<210> 22

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: synthetic DNA

<400> 22

ggcttgggca cggcctga

18

<210> 23

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic DNA

<400> 23

gcggtgaaga ccaagctcaa actcactcca

30

<210> 24

<211> 21

<212> DNA

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<223> Description of Artificial Sequence: synthetic DNA

<400> 24

agaacctgcg tgcaatccat c

21

<210> 25

<211> 23

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic DNA

<400> 25

cccgatcatga gggcgtcggt ggc

23

<210> 26

<211> 27

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic DNA

<400> 26

accagcaacg gtgggcggtt ggtaatc

27

<210> 27

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic DNA

<400> 27

ggcacgcgac acgctgtg

18

<210> 28

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: synthetic DNA

<400> 28

agctagccgt gactagggct aagatggagc

30

<210> 29

<211> 20

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:synthetic
DNA(primer)

<400> 29

aacaagatgg attgcacgca

20

<210> 30

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:synthetic
DNA(primer)

<400> 30

cgtcaagaag gcgatagaag

20

<210> 31

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:synthetic DNA

<400> 31

gcactctctg cagtcattgcg gctcacggac

30

<210> 32

<211> 28

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:synthetic DNA

<400> 32

cccctgtgag gaactactgt cttcacgc

28

<210> 33

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence::synthetic DNA

<400> 33

ccgggagagc catagtggtc tgcg

24

<210> 34

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence::synthetic DNA

<400> 34

ccactcaaag aaaaagtgac acgagctcgc

30

<210> 35

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:synthetic
DNA(primer)

<400> 35

ggcttgggca cggcctga

18

<210> 36

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence::synthetic DNA

<400> 36

gcggtgaaga ccaagctcaa actcactcca

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<210> 37

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence::synthetic DNA

<400> 37

agaacctgcg tgcaatccat c

21

<210> 38

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence::synthetic DNA

<400> 38

cccgtcatga gggcgtcggg ggc

23

<210> 39

<211> 27

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence::synthetic DNA

<400> 39

accagcaacg gtgggcgggt ggtaatc

27

<210> 40

<211> 18

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<213> Artificial Sequence

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<223> Description of Artificial Sequence::synthetic DNA

<400> 40

ggaacgcgac acgctgtg

18

<210> 41

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence::synthetic DNA

<400> 41

agctagccgt gactagggt aagatggagc

30